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H0616: 3, L0665: 3, L0794: 2, L0659: 2, L0779: 2, H0423: 2,	H0171: 1, H0685: 1, H0675: 1, S0408: 1, H0013: 1-1-0105: 1	HO581: 1, T0110: 1, S0214: 1, H0622: 1	H0561: 1, S0002: 1,	H0529; 1, L0762; 1, L0667: 1, L0766: 1,	L0805: 1, L0655: 1,	L0658: 1, L0790: 1,	H0702: 1, H0648: 1,	H0521: 1, L0748: 1,	L0777: 1, L0758: 1 and	L0366: 1.				AR089: 18, AR060:	9, AR055: 6, AR061:	4	L0754: 12, L0740: 8,	1 0649. 6 HO620. 5	S0214: 5, L0766: 5,
									•			Asp-30 to Thr-37, Lys-81 to Glu-87.	Ile-13 to Gly-18.	Val-23 to Lys-30, Tvr-40 to Asn-50	Lys-63 to Lys-68,	Leu-92 to Asn-102.			
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	Asn-14 to Thr-26.			Leu-3 to Met-9, Gln-24 to Thr-32	Ser-25 to Cvs-36	Gln-39 to Asp-56.	Asn-49 to Asp-54.					-			•	•		
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	Val-77 to Val-86.	Pro-69 to Leu-76.																			
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Arg-5 to Arg-17, Asp-25 to Val-31.	Val-7 to Glu-13,	Thr-20 to Arg-26,	Lys-34 to Phe-40,	Arg-46 to Lys-51.	Lys-27 to Gln-39,	Gln-47 to Thr-52.	Leu-3 to Glu-14,	Ser-95 to Ala-105.		Pro-24 to Val-49,	Tyr-51 to Leu-56.															-		
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		Leu-31 to Met-36, H06 Pro-97 to Lys-103, L07 Gln-126 to Trn-136 H00
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Pro-11 to His-21,	Glu-30 to Glu-41,	Arg-55 to Ser-64,	Arg-89 to Thr-96,	Cys-99 to Gln-106,	Pro-116 to His-124.							-							-										
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HO	20)H	OH HO	<u> </u>	<u>80</u>	OH H)H)H	HC	HC)HI)H	<u>2</u>	<u>유</u>	<u>80</u>	<u> </u>	H)H	0S)H	요.)H	os	<u> </u>	<u> </u>	<u>.</u>)H)H	HC
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	Glu-1 to Gln-8, Pro-31 to Trp-36.																				
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H0553: 2, H0644: 2,			
H0271: 2, H0030: 2,			
H0375: 2, H0266: 2,	-		,
H0024: 2, T0010: 2,			
H0046: 2, H0050: 2,			
H0597: 2, H0546: 2,			
H0156: 2, H0263: 2,		 	
S0222: 2, H0427: 2,		 ······································	
S0132: 2, H0549: 2,			
S0045: 2, S0046: 2,		 	
 H0295: 2, H0650: 2,			
H0686: 2, S0040: 2,			
H0352: 3, L0353: 2,			
L0759: 3, L0591: 3,			
S0122: 3, H0682: 3,			
L0663: 3, S0126: 3,	•		
L0783: 3, L0809: 3,		 	
L0774: 3, L0657: 3,			
L0521: 3, L0768: 3,			
L0769: 3, L0667: 3,		 	
L0564: 3, S0372: 3,			
H0031: 3, H0264: 3,		 	
 S0388: 3, T0023: 3,		 	
S0010: 3, H0596: 3,			
H0431: 3, H0370: 3,		 	
L0603: 4, H0556: 3,			

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SC	<u> </u>	<u> </u>	<u> </u>	3	77	<u>고</u>	H	H	H	<u>80</u>	H	H	<u>83</u>	<u>3</u>	3	3	H	H	H	H	H	<u>80</u>	H	H)H	ដ	08)H	HO
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1, H0550: 1	, H0587: 1	I, H0331:	H0574: 1, H0559: 1	F0039: 1, H0013: 1	H000	T00	S0474: 1, H0581:	H02	H02	, H0051: 1	H05	1, S0334: 1,	S0250:	l, H0119:	H06	H0617: 1, L0055: 1	H01	H05	JO T	7,	H0560: 1, H0561: 1	S0352: 1, S0438: 1	H01:	H0641: 1, H0647: 1	H0646: 1, S0344:	S0426: 1, L0598: 1	107) 	.0771: 1, L0773: 1
59: 1,		33: 1,	74: 1,	39: 1,	30: 1,	42: 1,	74: 1,	9.1,	99: 1,	H0011: 1,	H0021: 1,	H0594: 1,	H0687: 1,	4: 1,	6: 1,	17: 1,	32: 1,	24: 1,	51: 1,	28: 1,	60: 1,	52: 1,	5.1,	41: 1	46: 1	26: 1,	29: 1	46: 1,	71: 1,
H0369: 1	H0455:	H0333: 1	H05	<u> 1003</u>	S028	700H	S047	S0049: 1	H0309:	00H	H00.	H05	H06	S0214: 1	<u>100</u>	90H	H00.	H01.	H05	H01;	H05	S035	S 04)H06	90H	S042	H05	106	107
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		Pro-1 to Arg-6, Met-12 to Ser-21.			Ser-35 to Ser-41,	Arg-1 to Gly-7.	
	3200	3201	3202	3203	3204	3205	3206
	33 - 164	3 - 398	3-392	15 - 455	1 - 201	376 - 564	113 - 475
	1025	1026	1027	1028	1029	1030	
	HAOTH28 HAOTH28R	HAOTI07 HAOTI07R	HAOTT79R	HAOTU79 HAOTU79R	HAOTW22R	HAOTW30 HAOTW30R	HAOTW65 HAOTW65R
	HAOTH28	HAOTI07	HAOTT79	HAOTU79	HAOTW22 HAOTW22	HAOTW30	HAOTW65

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	Lys-1 to Ser-6,	Ala-26 to Thr-31.	Ser-1 to Pro-11.				Ser-13 to Arg-19, Ile-28 to Asn-33.													-		Ala-19 to Arg-24.	Gly-1 to Thr-12,	Glu-23 to Lys-38.	Ser-119 to Lys-124.	Lys-32 to Arg-40,
3207	3208		3209	3210	3211	3212	3213	3214														3215	3216		3217	3218
68 - 232	200 - 313		2 - 193	1 - 258	208 - 396	3 - 299	84 - 203	3-116							-							1 - 165	108 - 239		3 - 587	2 - 517
1032	1033		1034	1035	1036	1037	1038	1039														1040	1041		1042	1043
HAPNK45R	HAPPR43R		HAPQO14 HAPQO14R	HAOML40 HAOML40R	HASME85R	HAUAK54R	HAVMB46 HAVMB46R	HAVMW69 HAVMW69R														HAVNQ92R A	HAVOJIIR	-	HAZAC68R	HAZAD13R
HAPNK45	HAPPR43		HAPQ014	HAQML40	HASME85	HAUAK54	HAVMB46	HAVMW69														HAVNQ92 HAVNQ92R A	HAVOJ11		HAZAC68	HAZAD13

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Ser-54 to Gly-59, Glu-133 to Ala-147, Gln-153 to Gln-164.	Val-9 to Gly-24.		Gln-12 to Thr-20,	Cys-39 to Asn-45,	Thr-60 to Gly-66,	Lys-99 to Asp-106,	Asp-119 to Pro-127.															
	3219	3220	3221																	•		
	191 - 445	3 - 512	2 - 466																			
	1044	1045	1046																			
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99: 4.	585: 3,	83: 3,	186: 3,	309: 3,	517: 3,	4:3,	67:3,	94: 3,	06: 3,	28: 3,	26: 3,	558: 3,	30: 3,	28: 3,	39: 3,	57: 3,	49: 2,	41:2,	HO2: 2,	22: 2,	92: 2,	01: 2,	121: 2,	31:2,	23: 2,	71:2,	4.2, 2,	24: 2,	16: 2,
4, L ₀₅	4, HO	3, H04	3, H ₀ /	3, HO	3, H00	3, S01	3, L07	3, L07	3, 1.08	3, L05	3, S01	3, H06	3, S03	3, 500	3,1,04	3, L07	3, H01	2, H03	2, H04	2, H07	2, H03	2, H01	2, H04	2, H02	2, H00	2, H02	2, H06	2, H01	2, H06
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59: 2	509: 2	22: 2,	2:2	51:2	589: 2	559: 2	548: 2	4.2.2	43: 2	88: 2	542: 2	300: 2	524: 1	171:1	24: 1	294: 1	557: 1	760: 1	255: 1	118:1	576: 1	08: 1	730: 1	78: 1	22: 1	330: 1	586: 1	522: 1	427: 1
Ĭ E	HOH	S0 40	<u>5</u>	<u>2</u>)H	<u>Ř</u>	HOH	<u>8</u>	5	<u>5</u>)H	<u> </u>	Ě	H	S60	HO	Ĕ	<u>5</u>	H0.	S04	<u>H</u>	<u>S</u>	H	<u>S</u> 02	802	H	HOH	ទ្ធ	HÓ
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	Pro-48 to Gly-59.	Gly-2 to Ser-9, Pro-46 to Cys-60, Pro-62 to His-69,	Pro-81 to Thr-91, Ala-96 to Ser-101.	Lys-11 to Ser-16, Arg-20 to Glu-28.
	3222	3223		3224
	282 - 467	138 - 515		1 - 165
	1047	1048		1049
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J0742: 8, S0410: 6,	.0756: 6, L0752: 6,	.0758: 6, L0731: 5,	H0341: 4, L0766: 4,	0803: 4, L0776: 4,	S0126: 4, H0170: 3,	H0486: 3, S0346: 3,	H0204: 3, H0545: 3,	.0471: 3, S0440: 3,	.0769: 3, L0772: 3,	.0804: 3, H0214: 3,	30404: 3, L0744: 3,	L0754: 3, L0757: 3,	H0506: 3, H0171: 2,	H0685: 2, S0420: 2,	S0356: 2, S0358: 2,	S0045: 2, H0587: 2,	H0069: 2, L0021: 2,	H0575: 2, T0048: 2,	H0318: 2, H0581: 2,	H0597: 2, H0009: 2,	H0014: 2, H0510: 2,	H0622: 2, H0031: 2,	S0364: 2, H0591: 2,	H0056: 2, H0494: 2,	S0438: 2, L0770: 2,	L0761: 2, L0521: 2,	L0649: 2, L0650: 2,	L0526: 2, L0792: 2,	.0666: 2, L0665: 2,
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H0413: 1, T0069: 1, S0038: 1, T0042: 1, H0512: 1, H0560: 1, H0646: 1, S0344: 1, S0422: 1, L0763: 1, L0764: 1, L0773: 1, L0774: 1, L0773: 1, L0774: 1, L0773: 1, L0659: 1, L0659: 1, L0659: 1, L0779: 1, L0789:	L0659: 6, L0662: 5, S0212: 4, S0003: 4, H0551: 4, S0192: 4,
	Phe-34 to Lys-41.
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S0376: 3, S0440: 3,	S0374: 3, H0521: 3,	S0040: 2, S0418: 2,	S0444: 2, H0580: 2,	10486: 2, H0575: 2,	,0471: 2, H0024: 2,	H0553: 2, L0637: 2,	126: 2, S0406: 2,	S0028: 2, L0748: 2,	754: 2, L0757: 2,	S0434: 2, L0590: 2,	242: 2, H0686: 1,	H0685: 1, H0664: 1,	420: 1, L0005: 1,	S0360: 1, S0408: 1,	045: 1, S0476: 1,	S0278: 1, H0357: 1,	H0586: 1, T0039: 1,	H0635: 1, H0427: 1,	.0022: 1, H0004: 1,	H0505: 1, S0049: 1,	H0251: 1, H0263: 1,		H0051: 1, H0356: 1,		H0090: 1, H0412: 1,	H0623: 1, H0560: 1,	H0641: 1, H0633: 1,	S0142: 1, S0208: 1,	.0773: 1, L0521: 1,
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L0649: 1, L0803: 1, L0775: 1, L0528: 1, L0663: 1, L0664: 1, H0144: 1, L0565: 1, H0693: 1, H0689: 1, H0690: 1, H0689: 1, H0659: 1, H0658: 1, H0670: 1, H0660: 1, S0152: 1, H0555: 1, H0436: 1, S0432: 1, L0439: 1, L0758: 1, L0686: 1, S0436: 1, L0688: 1, S0436: 1, L0688: 1, S0196: 1,		S0420: 2, H0575: 2, S0418: 1, L0005: 1, S0356: 1, H0619: 1, H0097: 1, H0581: 1, H0266: 1, S0250: 1, T0042: 1, H0625: 1, H0561: 1, S0438: 1, S0150: 1, H0691: 1, S0210: 1, H0690: 1, H0519: 1, H0690: 1,
	Ser-54 to Ser-60, Ile-69 to Gly-74.	Ala-17 to Asp-36, Thr-56 to Met-61.
·	3226	3227
	3 - 260	1 - 612
	1051	1052
	HAZAI89R	HAZAJ72R
	HAZAI89	HAZAJ72

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		L0748: 31, L0747: 18,	L0752: 17, L0758: 17,	L0777: 15, L0766: 13,	H0014: 9, L0776: 9,	L0751: 9, L0740: 7,	L0779: 7, L0754: 6,	H0040: 5, L0769: 5,	S0330: 5, L0439: 5,	L0756: 5, L0599: 5,	S0356: 4, H0587: 4,	H0622: 4, L0783: 4,	L0744: 4, L0745: 4,	L0749: 4, L0755: 4,	S0031: 4, H0624: 3,	S0222: 3, H0156: 3,	S0010: 3, S0049: 3,	S0051: 3, S0003: 3,	H0591: 3, H0038: 3,	L0763: 3, L0770: 3,	L0662: 3, L0551: 3,	L0803: 3, L0664: 3,	H0144: 3, H0547: 3,	H0521: 3, L0757: 3,	L0759: 3, L0596: 3,	L0608: 3, L0595: 3,	S0114: 2, S0116: 2,	H0341: 2, S0418: 2,
Thr-41 to Lys-46, Thr-53 to Gly-59,	Ala-77 to Asp-84.	Ser-1 to Leu-14,	Glu-39 to Thr-47.																	-						<u>-r - 1,</u>		
3228	•	3229											_															
27 - 389	-	2 - 172	٠												-			•										
1053		1054										•	·															
HAZAQ80R		HAZAW22 HAZAW22R					•																					
HAZAQ80		HAZAW22																										

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H0619	, H0013	, H0318	, H0051	H0428	, H0673	, H0674	H0616	, H0268	,1.0768	,10775	, L0805	, L0731	H0445	L0605	, H0543	, H0171	, S6024	, S0218	, H065(, L0785	, S0282	, \$0030	, H063	, \$0420	, S0442	, \$0476	, \$0300	, S0278: 1	, H043
S0360: 2, H0619: 2,	H0393: 2, H0013: 2,	H0575: 2, H0318: 2,	H0581: 2, H0051: 2,	S0388: 2, H0428: 2,	70006: 2, H0673: 2,	H0169: 2, H0674: 2,	S0036: 2, H0616: 2,	H0551: 2, H0268: 2,	.0625: 2, L0768: 2,	.0649; 2, L0775; 2,	0806: 2	H0539: 2, L0731: 2,	S0260: 2, H0445: 2,	30434: 2, L0605: 2,	0594: 2	H0170: 1, H0171: 1	H0716: 1, S6024: 1,	0134: 1	H0583: 1, H0650: 1	10657: 1	0212: 1	S0035: 1, S0030: 1,	H0305: 1, H0638: 1	S0348: 1, S0420: 1	.0005: 1, S0442: 1,	H0340: 1, S0476: 1,	10645: 1	H0411: 1,	H0549: 1, H0431: 1
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H0409: 1, H0331: 1,	H0574: 1, H0486: 1,	S0346:_1, H0421: 1,	052: 1, H0251: 1,	032: 1, H0085: 1,	H0545: 1, H0457: 1,	041: 1, H0050: 1,	H0024: 1, H0373: 1,)28: 1, S0250: 1,	615: 1, H0688: 1,	039: 1, S0364: 1,	H0361: 1, L0455: 1,	H0376: 1, H0135: 1,	090: 1, H0634: 1,	F0004: 1, T0042: 1,	S0014: 1, H0560: 1,	H0625: 1, S0438: 1,	H0509: 1, H0646: 1,	H0529: 1, L0637: 1,	L5575: 1, L0667: 1,	546: 1, L0764: 1,	771: 1, L5574: 1,	774: 1, L0655: 1,	527: 1, L0657: 1,	559: 1, L0809: 1,	789: 1, L0792: 1,	L0793: 1, L0663: 1,	.0665: 1, S0053: 1,	H0520: 1, H0519: 1,	S0126: 1, H0711: 1,
0H]	OH	203	OH .	<u></u>	HO	707)OH)98	OH.	OH	OH	H0.	OH.		200	OH	HO	OH	1.5	<u> </u>		101	<u>3</u>	Local Control of the	<u> </u>	207	<u> </u>	HO	S01
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H0690: 1, H0660: 1, S0136: 1, H0522: 1, S0406: 1, H0522: 1, S3012: 1, S0028: 1, L0743: 1, L0750: 1, L0750: 1, L0361: 1, S0026: 1, H0653: 1, S0196: 1, H06423: 1, S0412: 1 and S0424: 1	S0003: 154, S0214: 41, H0545: 38, L0651: 34, H0546: 17, L0747: 15, L0757: 14, L0731: 10, L0749: 6, H0544: 5, S0122: 5, H0365: 5, L0740: 5, S0360: 4, L0771: 4, L0662: 4, L0771: 4, L0662: 4, L0779: 3, H0169: 3, L0760: 2, L0764: 2, L0766: 2, L0764: 2, L0766: 2, L0764: 2, L0766: 2, L0765: 2, L0766: 2, L0759: 2, L0766: 2, L0759: 2, L0754: 2, L0756: 2, L0754: 2, L0756:
	3230
	225 - 359
	1055
	HAZBH84R
	HAZBH84

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H0411: 1, H0357: 1, H0486: 1, L0021: 1, H0309: 1, H0231: 1, L0738: 1, H0039: 1, S0366: 1, H0038: 1, S0142: 1, S0002: 1, L0762: 1, L0640: 1, L0763: 1, L0640: 1, L0523: 1, L0657: 1, L0523: 1, L0655: 1, L0555: 1, L0664: 1, L0565: 1, S0008: 1, H0690: 1, S0328: 1, H0690: 1, S0328: 1, L0758: 1, H0595: 1 and	H0506: 1. L0758: 3, H0068: 2, L0749: 2, L0756: 2, L0731: 2, S0026: 2, H0136: 2, S0040: 1, L0005: 1, H0351: 1, H0318: 1, H0052: 1, H0530: 1, H0412: 1, S0003: 1, H0412: 1, S0003: 1, H0529: 1, H0517: 1, L0762: 1, L0763: 1, L0770: 1,
	Ser-29 to Ile-39, Gly-42 to Trp-48, Lys-82 to Val-92.
	3231
	56 - 424
	1056
	HAZBI39R
	HAZBI39

L0769: 1, L0766: 1, L0775: 1, L0809: 1, H0698: 1, H0690: 1, S0390: 1, S0028: 1, L0742: 1, L0748: 1, L0775: 1, L0779: 1, L0759: 1, S0308: 1, H0542: 1, H0543: 1 and H0423: 1.					L0665: 3, H0318: 2,	L0659: 2, S0330: 2,	L0439: 2, L0599: 2,	H0583: 1, H0038: 1,	L0761: 1, L0771: 1,	L0387: 1, L0655: 1,	L0666: 1, L0779: 1,	L0777: 1, L0758: 1 and	L0608: 1.		AR052: 12, AR055: 8,	8, AR033:	AR089: 6, AR061: 6,	ARU60: 5, ARU96: 5,
	Gly-1 to Pro-6, Lys-58 to Asn-71.	Pro-15 to Thr-22, Lys-28 to Arg-35.												Asn-30 to Ile-35.	Gly-1 to Pro-38,	Leu-45 to Gly-64.		
	3232	3233	3234	3235	3236									3237	3238			
	3 - 323	207 - 371	2 - 199	3 - 245	267 - 446									94 - 261	1 - 297			
·	1057	1058	1059	1060	1061									1062	1063			
	HAZBJ69R	HBAGS73R	HBBBE13R	HBCPI58RA	HBJAC26R									HBJFS85R	囯			
	HAZBJ69	HBAGS73	HBBBE13	HBCPI58	HBJAC26									HBJFS85	Γ.			

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1, AR039:	L0794: 5, L0766: 3,	12: 2,	L0803: 2,	.0378: 2,	.0438: 2,	.0749: 2,	L0599: 2,	86: 1,	34: 1,	69: 1,	70: 1,	46: 1,	50: 1,	,66: 1,	83: 1,	63: 1,	16: 1,	26: 1,	69: 1,	46: 1,	58: 1,	75: 1,	59: 1,	09: 1,	91: 1,	59: 1,	52: 1,	7: 1,	77: 1,
	: 5, LO	3, S02	2, L08		_	$\overline{}$	_	2, H06	1, S013	1, H06	1, H03	1, S03	1, H00	1, H0266: 1	1, L04	1, H0163: 1	1, H0116:	•	1, L0769: 1			1, L0375:	1, L0659:	1, L0809:	1, H0691:	-	1, S0152:	1, 5002	1, 1.077
AR104:	L0794	.0759: 3, S0212: 2,	H0318: 2, 1	0804: 2,	.0665: 2,	.0439: 2,	.0591: 2,	.0362: 2, H0686: 1	S0342: 1, S0134:	H0656: 1, H0669:	S0358: 1, H0370:	H0069: 1, S0346:	H0251: 1, H0050:	H0373:	S0022: 1, L0483:	H0553:	.0600E	H0509: 1	H0529: 1,	.0638:]	.0648:]	.0381: 1	.0653: 1	.0782: 1,	.0793: 1,	H0435: 1	H0648: 1	\$0390: 1, \$0027:	S0028: 1, L0777:
4	•	<u> </u>	<u> </u>	브		브	<u> </u>	<u> </u>	S	<u>.11</u>	S	<u> </u>	بلز	<u>, 141</u>	S	<u>iLi</u>	<u>;11;</u>	<u>;11</u>	<u>111</u>	<u> </u>	<u> </u>		브	브		프	Ŧ	<u>(V)</u>	S
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L0731: 1, S0260: 1, L0596: 1, S0192: 1 and S0424: 1.		S0280: 1 and S0038: 1.											H0651: 9, L0601: 5,	H0559: 4, H0494: 4,	L0646: 3, H0484: 2,	H0255: 2, S0358: 2,	H0550: 2, H0309: 2,	H0014: 2, H0188: 2,	H0617: 2, H0646: 2,	L0769: 2, L0644: 2,	L0806: 2, H0689: 2,	L0750: 2, L0603: 2,	H0624: 1, H0171: 1,	H0657: 1, H0663: 1,
	Glu-8 to Asn-13, Arg-16 to Thr-29.	Lys-23 to Lys-32, Ser-72 to Leu-80,	Ser-90 to Gly-95,	Leu-98 to His-103,	Ser-106 to Gly-113,	Leu-2 to Val-10.	His-2 to Glu-28,	Ala-37 to Gly-45,	His-54 to Pro-62,	Glu-69 to Thr-77.	Arg-1 to Arg-6,	Phe-27 to Arg-32.												
	3239	3240		•		3241	3242				3243		3244	!										
·	145 - 231	59 - 547				1 - 126	145 - 543				171 - 359		2 - 322											
	1064	1065				1066					1068		1069											
	HBJMN13R	HBXCG52R				HRXFO08R	HCACS53 HCACS53RP	400			HCHAJ85R		UCUMANTI HCHWM71R								-			
	HBJMN13	HBXCG52				HRYFOOR	HCACS53				HCHAJ85		וכשעשטח	nCnivily / 1										

S0360: 1, H0574: 1, H0257: 1, H0013: 1, H0635: 1, H0098: 1, H0253: 1, H0327: 1, H0545: 1, H0024: 1, H0181: 1, H0359: 1, S0150: 1, L0369: 1, L0771: 1, L0662: 1, L0766: 1, L0803: 1, L0661: 1, S0428: 1, H0670: 1, S0328: 1, L0748: 1, L0439: 1, L0758: 1 and H0543: 1.			L0439: 8, H0052: 7, L0748: 7, S0440: 5, L0758: 5, H0046: 4, H0038: 4, L0769: 4, S0442: 3, H0013: 3, H0253: 3, T0010: 3, L0774: 3, L0776: 3, H0144: 3, L0776: 3, L0731: 3, H0551: 3, S0404: 3, L0752: 3, L0731: 3, H0656: 2, S0358: 2, S0360: 2, S0222: 2, H0618: 2, H0620: 2, L0351: 2,
	Ser-1 to Thr-8, Glu-17 to Ala-32, Arg-39 to Trp-47.		Gln-17 to Pro-27.
	3245	3246	3247
	53 - 319	3 - 134	234 - 443
	1070	1071	1072
	HCHMW20 HCHMW20R	HCLBH21R	HCNDT49R
·	HCHMW20	HCLBH21	HCNDT49

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S0422: 2, L0764: 2,	L0771: 2, L0783: 2,	L0793: 2, H0658: 2,	H0666: 2, L0751: 2,	L0754: 2, L0745: 2,	L0747: 2, L0750: 2,	H0624: 1, H0265: 1,	H0556: 1, H0686: 1,	S0134: 1, S0212: 1,	S0001: 1, H0254: 1,	H0661: 1, S0354: 1,	S0444: 1, S0408: 1,	S0300: 1, S0278: 1,	H0369: 1, H0370: 1,	H0333: 1, H0574: 1,	H0486: 1, H0036: 1,	H0263: 1, H0597: 1,	H0545: 1, H0572: 1,	H0024: 1, S0388: 1,	S0051: 1, S0250: 1,	H0252: 1, H0428: 1,	H0039: 1, H0644: 1,	L0055: 1, H0674: 1,	H0135: 1, H0087: 1,	F0067: 1, H0488: 1,	H0529: 1, L0763: 1,	L0770: 1, L0761: 1,	L0374: 1, L0662: 1,	L0768: 1, L0766: 1,	L0803: 1, L0775: 1,
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L0805: 1, L0653: 1, L0526: 1, L5622: 1, L0666: 1, L0664: 1, L0665: 1, S0053: 1, H0547: 1, H0682: 1, H0660: 1, H0670: 1, H0660: 1, H0670: 1, H0672: 1, S0328: 1, H0672: 1, S0328: 1, H0696: 1, S0048: 1, S0406: 1, H0631: 1, S0406: 1, L0749: 1, L0742: 1, L0749: 1, L0759: 1, S0436: 1, S0011: 1, S0192: 1, H0542: 1, H0423: 1,		L0747: 15, L0750: 7, L0731: 7, L0588: 6, L0748: 5, L0439: 5, L0759: 5, H0150: 3, S0144: 3, L0666: 3, L0665: 3, L0752: 3, L0758: 3, L0591: 3, S6024: 2, T0049: 2, S0356: 2, H0674: 2, H0413: 2, S0142: 2,
	Arg-1 to Leu-10, Ser-32 to Ala-45, Ala-58 to Gly-68.	Tyr-3 to Gly-10, Glu-19 to Trp-25.
·	3248	3249
·	25 - 273	32 - 754
	1073	1074
	HCOMA72 HCOMA72R	HCOMB04R
	HCOMA72	HCOMB04

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.0770: 2, L0769: 2,	.0774: 2, L0775: 2,	.0657: 2, H0672: 2,	30328: 2, S0378: 2,	.0605: 2, L0593: 2,	H0624: 1, T0002: 1,	10638: 1, S0418: 1,	L0005: 1, S0376: 1,	30444: 1, S0360: 1,	H0675: 1, S0045: 1,	H0393: 1, S0222: 1,	H0441: 1, H0453: 1,	H0333: 1, H0574: 1,	H0486: 1, H0101: 1,	L0021: 1, H0098: 1,	r0010: 1, H0083: 1,	H0266: 1, H0188: 1,	S0003: 1, H0328: 1,	H0615: 1, H0428: 1,	T0006: 1, H0166: 1,	H0598: 1, H0090: 1,	H0063: 1, T0069: 1,	H0509: 1, H0641: 1,	S0002: 1, L0369: 1,	L0372: 1, L0662: 1,	L0794: 1, L0766: 1,	L0803: 1, L0651: 1,	L0655: 1, L0379: 1,	L0656: 1, L0659: 1,	L0782: 1, L0783: 1,
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L0788: 1, L0664: 1, L0565: 1, L0438: 1, H0547: 1, S0126: 1	H0435: 1, H0659: 1, H0658: 1, H0670: 1,	H0522: 1, H0696: 1,	L0740: 1, L0751: 1, L0745: 1, L0755: 1,	L0757: 1, H0445: 1,	H0343: 1, H0595: 1,	L0596: 1, L0581: 1, L0599: 1 and S0192: 1.			L0740: 6, L0666: 2,	H0547: 2, L0751: 2,	L0777: 2, H0483: 1,	S0358: 1, S0444: 1,	H0038: 1, L0770: 1,	•	- :	L0789: 1, H0520: 1,	H0659: 1, H0670: 1,	L0747: 1, L0779: 1 and	L0758: 1.	L0439: 7, L0748: 5,	H0255: 4, H0253: 4,	H0012: 3, H0620: 3,	H0617: 3, L0783: 3,
				-			Pro-1 to Pro-9, Ile-43 to Lys-48.		Asn-10 to Val-16.											Lys-11 to Thr-16,	Arg-18 to Glu-24,	Arg-31 to Leu-38,	Phe-51 to Glu-56,
							3250	3251	3252		_									3253			
							3 - 275	1 - 315	350 - 622											2 - 523			
						_	1075	1076	1077											1078	•		
·							HCOMD38R	HCOMD61R	HCOMF10R											HCOMF52R			
							HCOMD38	HCOMD61	HCOMF10 HCOMF10											HCOMF52			

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L0438: 3, H0618: 2,	H0046: 2, L0769: 2,	L0766: 2, L0532: 2,	L0663: 2, H0521: 2,	L0750: 2, L0777: 2,	L0592: 2, H0543: 2,	H0650: 1, S0212: 1,	S0001: 1, H0483: 1,	H0254: 1, H0638: 1,	S0418: 1, S0358: 1,	H0580: 1, S0468: 1,	H0549: 1, H0559: 1,	H0013: 1, S0280: 1,	H0318: 1, H0052: 1,	H0024: 1, H0252: 1,	H0424: 1, H0553: 1,	H0644: 1, H0124: 1,	H0376: 1, H0135: 1,	H0040: 1, H0087: 1,	H0488: 1, T0042: 1,	H0494: 1, H0509: 1,	L0768: 1, L0388: 1,	L0657: 1, L0659: 1,	L0789: 1, L0352: 1,	S0126: 1, H0689: 1,	H0658: 1, H0670: 1,	H0436: 1, H0478: 1,	H0626: 1, L0742: 1,	L0780: 1, L0731: 1,	H0445: 1, L0608: 1,
	Thr-102 to Thr-114,	Leu-119 to Gln-124.					<u> </u>																						
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L0366: 1 and S0276: 1.											H0670: 14, S0360: 9,	L0657: 5, L0659: 4,	L0809: 4. H0696: 4.	10777: 4 \$0356: 3	1.0764: 3, 1.0662: 3.	S0374: 3. H0637: 2.	U0411: 2 U0486: 2	HO411: 2, HO486: 2,	L0762: 2, L0763: 2,	L0521: 2, L0768: 2,	L0774: 2, L0805: 2,	L0666: 2, L0665: 2,	H0659: 2, H0672: 2,	H0710: 2, S0406: 2,	L0744: 2, L0748: 2.	H0686: 1, S0040: 1.	H0713: 1, H0657: 1,	H0662: 1, S0444: 1,	S0408: 1, S0410: 1,
	Lys-1 to Arg-8, Asp-20 to Thr-26,	Gly-35 to Glu-48,	Leu-66 to Arg-72,	Phe-83 to Tyr-88.	Trp-13 to Asp-19,	Cys-29 to Gln-34,	Ala-41 to Arg-52,	Gly-54 to Gln-59,	Arg-69 to Tyr-83.		Pro-13 to Pro-22.																		
	3254				3255					3256	3257																		
	46 - 357				29 - 373					233 - 436	21 - 167							•				·	-						
	6/01				1080					1081	1082	-																	
	HCOMG28 HCOMG28R				HCOMG40 HCOMG40R					HCOMG82R	HCOMH70 HCOMH70R																		
	HCOMG28				HCOMG40	٠				HCOMG82	HCOMH70								,				-						

Pu			: 13, 11,
H0602: 1, H0643: 1, H0596: 1, H0263: 1, H0015: 1, H0014: 1, H0015: 1, H0014: 1, H0015: 1, H0687: 1, S0003: 1, S0214: 1, H0169: 1, H0674: 1, L0640: 1, L0674: 1, L0640: 1, L0676: 1, L0640: 1, L0676: 1, L0648: 1, L0806: 1, L0784: 1, L0806: 1, L0776: 1, L0655: 1, L0776: 1, L0659: 1, H0547: 1, H0648: 1, S0328: 1, S0330: 1, H0555: 1, H0627: 1, S0328: 1, S0330: 1, L0731: 1, S0434: 1, L0731: 1, L0779: 1, L0731: 1, L0779: 1, L0731: 1, L0362: 1 and H0667: 1,			L0743: 19, L0731: 12 L0744: 12, L0748: 11, L0666: 9, L0751: 9,
:		Lys-1 to Gly-10,	Pro-13 to Asn-23.
	3258	3259	3261
	42 - 497	244 - 618 255 - 605	138 - 338
	1083	1084	
	HCOMI30R	HCOMI37R HCOML11R	HCOMM12 HCOMM12R
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.0664: 8, H0144: 8	, H0617:	.0769: 7, L0662: 7	H0616:	, L0483: 5,	, L0517:	.0519: 5, L0754: 5,	, \$0051:	L0663:	H0658: 4, L0747: 4	.0750: 4, L0757: 4,	H0295: 3, T0049: 3	, L0622: 3	10039: 3, H0253: 3	, L0163:	H0594: 3, H0188: 3	, H0100:	L0372:	, L0776: 3	, L0382: 3	, H0672: 3	L0742:	J0752: 3, S0260: 3,	, L0596: 3,	.0588: 3, L0599: 3,	H0637:	S0132:	S0278: 2,	H0431:	, H0599:
L0664: 8	Н0333: 7, Н0617: 7	L0769: 7	S0022: 6, H0616: 6,	H0087: 6, 1	H0181: 5, L0517: 5,	L0519: 5	H0677: 5, S0051: 4	S0344: 4, L0663: 4,	H0658: 4	L0750: 4	H0295: 3	H0255: 3, I	Г0039: 3	H0545: 3, L0163: 3	H0594: 3	H0673: 3, H0100: 3	S0142: 3, L0372: 3,	L0775: 3, I	L0657: 3, 1	L0665: 3,]	S0027: 3, L0742: 3,	L0752: 3	H0445: 3, I	L0588: 3	S0420: 2, H0637: 2,	S0045: 2, S0132: 2,	\$0300: 2,	S0222: 2, H0431: 2,	H0392: 2, H0599: 2
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-	S0134: 1, H0650: 1,		
	H0294: 1, S0114: 1,		
-	S0040: 1, S0342: 1,		
	T0002: 1, H0686: 1,		
	L0600: 2, H0556: 1,		
	S0192: 2, S0424: 2,		
	L0362: 2, L0601: 2,		
	L0740: 2, L0485: 2,		
	H0521: 2, S0028: 2,		
	H0670: 2, H0651: 2,		
	S0374: 2, S0126: 2,		
	L0809: 2, L0532: 2,		
	L0655: 2, L0384: 2,		
	L0804: 2, L0375: 2,		-
	L0771: 2, L0803: 2,		
	L0638: 2, L0764: 2,	-	
	L0763: 2, L0770: 2,		
	H0529: 2, L0369: 2,		
	H0038: 2, H0413: 2,		-
	S0036: 2, H0163: 2,		-
	H0428: 2, H0124: 2,		
	H0597: 2, H0544: 2,		
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	r0007: 1	S0007: 1	H0441:	H0587: 1,	L0623: 1	F0114: 1,	L0021: 1	H0004: 1	S0010: 1	S0049: 1	H0596:	H0327: 1	H0123:	H0006: 1,	H0051: 1	H0510: 1	H0687: 1	T0006: 1	H0182: 1,	H0383:	H0166:	H0135:	Γ0067: 1	H0412:	S0112: 1	H0494:	H0561: 1,	L0640: 1	.0761: 1
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Pro-10 to Leu-20, His-27 to Lys-40.	Phe-7 to Asp-29,	Glu-31 to Lys-38,	Thr-46 to Thr-52, Glu-65 to Arg-72.																						-
3262	3263			3264	3265	3266	3267																	3268	3269
298 - 119	1 - 336		,	3 - 443	166 - 492	3 - 161	76 - 267																	8 - 718	33 - 398
1087	1088			1089	1090	1001	1092																	1093	1094
HCOMM55R	HCOMOS8 HCOMOS8R			HCOMW52 HCOMW52R P00B	HCOMX77 HCOMX77R	HCONC18 HCONC18RP 00B	HCONH28R																	HCONK56R	HCONK75R
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	L0748: 50, L0754: 42,	L0731: 31, H0644: 30,	L0747: 29, H0551: 22,	S0418: 21, L0659: 17,	H0619: 15, L0663: 14,	L0666: 13, S0360: 10,	H0622: 10, H0031: 10,	H0553: 10, L0662: 10,	S0126: 10, L0757: 10,	L0603: 10, S0420: 8,	H0486: 8, H0013: 8,	H0050: 8, H0623: 8,	L0717: 7, H0144: 7,	L0770: 6, L0750: 6,	H0427: 5, H0024: 5,	H0328: 5, H0030: 5,	L0773: 5, L0665: 5,	L0749: 5, H0713: 4,	T0039: 4, S0280: 4,	H0012: 4, H0038: 4,	L0794: 4, L0774: 4,	L0775: 4, L0805: 4,	H0624: 3, H0170: 3,	H0171: 3, S0356: 3,	H0329: 3, S0046: 3,	H0331: 3, H0575: 3,	L0471: 3, H0620: 3,
His-23 to Lys-34, Arg-49 to Leu-57, Pro-59 to Ser-66.		•							•			•															,
3270	3271	•																									
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S0250: 3, H0032: 3,	H0591: 3, L0598: 3,	L0803: 3, L0653: 3,	L0657: 3, H0670: 3,	S0406: 3, H0555: 3,	L0780: 3, L0605: 3,	H0667: 3, H0717: 2,	T0049: 2, S0212: 2,	H0661: 2, S0376: 2,	S0007: 2, H0208: 2,	S0476: 2, H0645: 2,	H0411: 2, H0437: 2,	H0550: 2, H0586: 2,	H0333: 2, H0599: 2,	H0042: 2, H0545: 2,	H0123: 2, H0105: 2,	H0014: 2, H0051: 2,	H0375: 2, L0142: 2,	H0598: 2, H0135: 2,	H0040: 2, S0372: 2,	L0520: 2, L0640: 2,	1.0769: 2, 1.5566: 2,	L0772: 2, L0646: 2,	L0771: 2, L0363: 2,	L0768: 2, L0375: 2,	L0654: 2, L0776: 2,	L0809: 2, L0438: 2,	H0547: 2, H0519: 2,	L0602: 2, L0743: 2,	L0744: 2, L0439: 2,
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5: 8, H): 8, H	: 7, H	: 6, L(7.5, H	3: 5, H	: 5, L(: 5, H	: 4, S(2: 4, H	: 4, L	1,4:	t: 3, H	I: 3, S	9: 3, S	I: 3, H	1:3, H	3, H	1: 3, L	3, L	7: 3, H	5: 3, H); 3, L	7:3, H	. 2, S	1: 2, S	7: 2, H	5.2, H	1: 2, H	0: 2, H
H0486	H0050: 8, H0623: 8,	L0717	1.0770	H0427	H0328	1.0773	10749	T0035	H001;	L0794	1.0775	H062	H017	H032	H033	1047	S025(H059	1080	1065	S040	1078	H066	T004	990H	2000	S0476	H041	H055
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H0546	H0081	T0003	H0275	S0003:	H0111	S0364:	H0090	H0087	H0412	H0100	H0561	T0369	10800	1.0764	1.0804	10806	1.0526	L0788	10793	H0691)690H	H0684	H0672	H0518	S0146	S3014
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				Val-1 to Ile-10,	Met-32 to Asn-42.	Gly-7 to Gly-21,	Pro-145 to Arg-152,	Ala-178 to Lys-183.		Gly-18 to Gln-39,	Leu-49 to His-60,	Glu-67 to Asp-92.		Thr-76 to Arg-92,	Pro-113 to Ala-118.	Pro-15 to Tyr-21.	Asn-1 to Pro-10.						Ala-15 to Gln-25,	Gly-42 to Asp-51,	Gln-82 to Lys-87,	Lys-113 to Lys-118.	
				3276		3277			3278	3279			3280	3281		3282	3283	3284	3285	3286	3287	. 3288	3289				3290
				3 - 413		1 - 669			312 - 488	3 - 380			3 - 575	1 - 429		85 - 234	2 - 151	44 - 457	101 - 352	2 - 334	3 - 482	1 - 426	1 - 354				1115 209 - 625
				1101		1102			1103	1104			1105	1106	!	1107		1109		1111	1112	1113	1114				
				HCONP44R		HCONO67 HCONO67R	,		HCONO71R	HCONR31 HCONR31R			HCONU03R	_		HCONW74 HCONW74R	HCONW90 HCONW90R	HCONW95 HCONW95R	HCOOB60 HCOOB60R	HCOOG32 HCOOG32R	HCOOG37 HCOOG37R	HCOOH93R	HCOOI71R		-		HCOOI79R
	-			HCONP44		HCONO67			HCONO71	HCONR31)		HCONI 103	HCONW62		HCONW74	HCONW90	HCONW95	HCOOB60	HC00G32	HC00G37	нсоон93	HC00171)			HCOOI79

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Lys-71 to Glu-76, Ala-95 to Asp-100, Asp-113 to Ser-134.	Leu-33 to Lys-39.			
3291	3292	3293	3294	3295
2 - 694	2 - 661	2 - 166	2 - 229	2 - 439
1116	1117	1118	1119	1120
HCOOM18 HCOOM18R	HCOOM73R	HC00046R	HC00S39R	HCOOT43R
HCOOM18	HCOOM73 HCOOM73	HC00046	HCOOS39	HCOOT43

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445: 2,	604: 2,	362; 2,	543: 2,	134: 1,)656: 1,	1484: 1,	420: 1,	354: 1,	717: 1,	550: 1,	1, H0574: 1,	T0060: 1,	H0108: 1,	H0037: 1,	H0309: 1,	T0115: 1,	H0327: 1,	S0051: 1,	H0061: 1,	l, H0288: 1,	1, H0039: 1,	H0644: 1,	H0032: 1,	H0316: 1,	087: 1,	T0042: 1,	S0438: 1,	l, H0132: 1,)633: 1,
.0759: 2, H0445: 2,	.0605: 2, L0604: 2,	.0595: 2, L0362: 2,	S0194: 2, H0543: 2,	S0040: 1, S0134:	H0583: 1, H0656:	S0116: 1, H0484:	H0177: 1, S0420:	005: 1, S0	408: 1, LO	H0369: 1, H0550:	S0222: 1, HC	•	H0250: 1, H(H0575: 1, H(F0048: 1, HC		•	_	H0083: 1, H(S6028: 1, HC	H0292: 1, H(H0111: 1, H(H0383: 1, H(H0598: 1, H0087:	H0059: 1, TC	•	H0509: 1, H(H0641: 1, H0633:
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S0210: 1, H0529: 1, L0520: 1, L0763: 1, L0765: 1, L0763: 1, L0765: 1, L0773: 1, L0364: 1, L0766: 1, L0806: 1, L0665: 1, L0657: 1, L0518: 1, L0657: 1, L0518: 1, L0657: 1, L0518: 1, H0659: 1, H0649: 1, H0666: 1, H0649: 1, H0666: 1, H0648: 1, H0672: 1, S0013: 1, H0699: 1, S013: 1, L0699: 1, S013: 1, L0699: 1, L0744: 1, S0206: 1, L0697: 1, S0242: 1, L0697: 1, S0242: 1, L0697: 1, S0242: 1, L0697: 1,			
8333333333EEEEE3001000		Ala-20 to Asp-25, Asp-38 to Ser-59, Asn-103 to Glu-109, Pro-111 to Ser-119.	
	3297	3298	3299
	3 - 257	58 - 534	153 - 311
	1122	1123	1124
	HCOOTISE HCOOUSER	HCOOW72 HCOOW72R	HCOOX01R A
	HC001156	HCOOW72	HCOOX01 HCOOX01

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L0751: 14, L0659: 11,	H0695: 10, L0666: 10,	.0757: 9, L0743: 8,	.0779: 8, H0581: 6,	.0769: 6, L0748: 6,	.0758: 6, H0556: 5,	10402: 5, S0360: 5,	H0014: 5, H0271: 5,	.0761: 5, L.0667: 5,	.0774: 5, L0653: 5,	S0406: 5, H0436: 5,	30358: 4, S0132: 4,	L0763: 4, L0776: 4,	.0809: 4, L0664: 4,	S0126: 4, L0731: 4,	S0280: 3, H0150: 3,	H0687: 3, L0662: 3,	J0775: 3, L0375: 3,	H0670: 3, L0749: 3,	S0434: 3, H0295: 2,	H0650: 2, S0408: 2,	H0580: 2, H0411: 2,	10586: 2, H0559: 2,	H0545: 2, H0050: 2,	S0051: 2, H0083: 2,	HO416; 2, H0688; 2,	H0617: 2, H0063: 2,	H0551: 2, S0344: 2,	.0643: 2, L0764: 2,	.0766: 2, L0649: 2,
3300 His-15 to Gly-29.	<u></u>		1		!		<u>. .</u>		ted.	<u>S</u>	<u>ss</u>		<u> </u>	<u>.</u>	<u>ss</u>	<u>14</u>		<u>щ</u>	S		<u>, 114</u>	14	<u>. 14 </u>	<u>S</u>	<u>.+-</u> i	<u>-144</u>	<u></u>		
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236 - 367																			•								,		
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		.0665: 2, S0428: 2,	S0332: 2, L0747: 2,	L0752: 2, H0624: 1,	H0171: 1, H0265: 1,	H0159: 1, H0713: 1,	S0218: 1, H0656: 1,	30116: 1, H0483: 1,	H0663: 1, H0638: 1,	S0356: 1, S0444: 1,	H0339: 1, S0046: 1,	S0476: 1, S0278: 1,	S6022: 1, H0550: 1,	S0414: 1, L0622: 1,	HO486: 1, H0635: 1,	H0427: 1, H0156: 1,	H0599: 1, H0052: 1,	H0309: 1, H0597: 1,	H0546: 1, L0041: 1,	H0123: 1, H0510: 1,	H0188: 1, T0006: 1,	.0055: 1, H0673: 1,	H0708: 1, H0708: 1,	H0316: 1, H0634: 1,	H0087: 1, H0059: 1,	S0352: 1, S0448: 1,	S0306: 1, L0640: 1,	.0371: 1, L0770: 1,	.0638: 1, L5565: 1,
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L0646: 1, L0642: 1, L0645: 1, L0642: 1, L0773: 1, L0648: 1, L0794: 1, L0805: 1, L0652: 1, L0661: 1, L0493: 1, L0638: 1, L0663: 1, S0374: 1, F0068: 1, H0689: 1, H0682: 1, H0659: 1, H0666: 1, H0679: 1, H0710: 1, H0539: 1, H0522: 1, H0696: 1, S0190: 1, S0404: 1, L0740: 1, L0744: 1, L0777: 1, L0759: 1, L0777: 1, L0759: 1, L0777: 1, L0361: 1 and S0194: 1,			L0748: 4, L0747: 4, L0779: 4, H0620: 3, L0770: 3, L0774: 3, L0622: 2, S0280: 2,
	Leu-13 to Gly-21, Ser-27 to Ser-42.	Asn-67 to Arg-73, Thr-96 to Gly-102.	Ala-1 to Leu-7, Gly-14 to Arg-19, Tyr-34 to Asp-41.
	3301	3302	3303
	203 - 349	1 - 315	3 - 446
	1126	1127	1128
	HCOOX48R	HCQOY43R	HCOPA60 HCOPA60RA
	HCOOX48 HCOOX48	HCOOY 43	HCOPA60

H0290: 2, 2, 20800: 2, 2, 20766: 2, 2, 20788: 2, 2, 20758: 2, 40295: 1, 40024: 1, 40024: 1, 40648: 1, 40648: 1, 40648: 1, 40653: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40652: 1, 40659: 1, 40652: 1, 40659: 1, 40652: 1, 40659: 1, 40669: 1, 4066	and S0196: 1.
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	Asp-28 to Glu-34, Ala-38 to Ile-44, Ala-64 to Arg-77, Gly-82 to Ala-88.
	3304
	2 - 598
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		L0751: 16, L0601: 10,	L0659: 4, H0670: 4,	L0665: 3, S0376: 2,	H0494: 2, L0758: 2,	L0596: 2, H0587: 1.	H0646: 1, L0372: 1.	L0646: 1, L0662: 1,	L0552: 1, L0376: 1,	L0806: 1, L0664: 1,	H0682: 1, H0435: 1,	HU6/2: 1 and LU/44: 1.						:	L0665: 3, H0682: 2,	H0435: 2, L0751: 2,	H0663: 1, L0021: 1,	H0647: 1, L0806: 1,	L0664: 1, H0683: 1,	H0670: 1 and H0660: 1.	-			
	Phe-1 to Gly-7.	Arg-37 to Gly-44,	Asp-76 to Gly-87.													Thr-6 to Cys-16,	Tyr-40 to Lys-47,	Ser-61 to Gln-66.	Arg-1 to Ser-7,	Ser-25 to Phe-32,	Leu-39 to Ser-44,		5.		Pro-7 to Phe-15,	Gly-26 to Arg-31,	Asp-35 to Glu-43.	Arg-4 to Lys-9,
3305	3306	3307			,							0000	2200	3309	3310	3311			3312						3313			3314
3 - 89	26 - 541	2 - 400										270	240 - 240	2 - 328	1 - 90	140 - 361			143 - 601						1 - 267			3 - 620
1130	1131	1132											1100	1134	1135	1136			1137						1138			1139
HCOPC45R	HCOPD67R	HCOPE27R										, d. Siraconi	HCOPH51KA	HCOPI09RA	HCOPM42R	HCOPO34R			HCOPO36R						HCOPO88R		•	HCOPV41R
HCOPC45	HCOPD67	HCOPE27						•						60IdODH	HCOPM42	HCOPO34			HCOPO36						HCOPO88			HCOPV41

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	AR096: 11, AR061: 10, AR104: 5, AR060: 5, AR089: 5, AR055: 4 L0774: 3, S0358: 2, H0509: 2, L0809: 2, H0670: 2, L0752: 2, H0449: 1, T0048: 1, H0355: 1, H0428: 1, H0529: 1, L0770: 1, L0761: 1, L0770: 1, L0761: 1, L0773: 1, L0803: 1, L0804: 1, L0375: 1, L0659: 1, L0375: 1, L0659: 1, L0382: 1, L0791: 1, S0328: 1, H0134: 1, L0743: 1, L0748: 1, L0747: 1 and L0756: 1.			-
Asp-61 to Asn-67, Thr-69 to Lys-75, Leu-105 to Ala-112, Gly-126 to His-134, Gly-172 to Glu-179, Pro-186 to Asn-199.	·	His-39 to Asn-45, Cys-68 to Cys-75.	His-37 to Asp-42.	Lys-6 to Asp-11, Gly-22 to Phe-27,
	3315	3316	3317	3318
	1 - 582	41 - 271	1 - 192	1 - 582
	1140	1141	1142	1143
	HCOPZ15R	HCOQA38 HCOQA38R	HCOQA80R	нсобв12 нсобв12к
	HCOPZIS	HCOQA38	HCOOA80	HCOQB12

	H0670: 2 and H0589:	,	L0794: 7, L0803: 5, 10809: 5, L0666: 5,	L0751: 5, L0759: 3, H0674: 2, L0768: 2,	.0774: 2, L0663: 2,	L0779: 2, L0758: 2,	30360: 1, H0455: 1,	H0618: 1, H0263: 1,	H0169: 1, H0616: 1,	H0059: 1, S0112: 1,	L0451: 1, L0770: 1,	.0639: 1, L0648: 1,		L0783: 1, L0789: 1,	20790: 1, L0791: 1,	L0664: 1, L0665: 1,	H0670: 1, H0672: 1 and	.0608: 1.		-		-			
His-54 to Gly-63, Asp-73 to Lys-79.		Lys-1 to Gln-7.	Lys-1 to Asp-21, Lys-61 to Leu-71.				<u> </u>		ш.				H					1	Glu-66 to Arg-72,	Pro-148 to Ser-154.	Met-3 to Ser-16,	Cys-25 to Asp-35,	Glu-42 to Lys-52,	Ser-55 to Glu-63,	Glu-74 to Thr-79,
	3319	3320	3321																3322	į	3323				
	1 - 66	281 - 466	3 - 248		-														1 - 570		106 - 639				
	1144	1145	1146																1147		1148				
	HCOQB56R	HCOQB58R	HCOQC71R																HCOQD29 HCOQD29R		HCOQD38R				
	нсоовзе	HCOQB58	нсоос71																HC0QD29		нсоор38				

	S0358: 15, S0003: 13, S0126: 11, L0754: 11, S0410: 10, H0599: 9, S0440: 9, H0529: 9, L0748: 9, H0529: 8, L0588: 8, L0362: 8, H0661: 7, S0007: 7, L0471: 7, L0731: 7, L0471: 7, L0731: 7, H0661: 7, S0007: 7, L0471: 7, L0731: 7, H0641: 7, H0644: 4, H0657: 4, S0346: 4, S0346: 4, S0346: 4, H0644: 4, H0644: 4, H0644: 4, H0644: 4, H0644: 4, L0750: 4, L0753: 4, L0750: 4, L0753: 4, L0750: 4, L0753: 4, L0758: 4, L0753: 4, L0758: 4, L0753: 3, H0663: 3, R0356: 3, S0356: 3, S0356: 3, S0356: 3, R0497: 3, S0322: 3, H0497: 3, S0322: 3, H
lle-81 to Ser-102, Ser-109 to Glu-118, Thr. 138 to Glu-158.	Asp-39 to Ser-46.
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36: 3,	52: 3,	47: 3,	80: 3,	39: 3,	95: 3,	14: 2,	12: 2,	125: 2,	76: 2,	80: 2,	69: 2,	109: 2,	574: 2,	56: 2.	118: 2,	96: 2,	63: 2,	.66: 2,	15: 2,	83: 2,	69: 2,	16: 2,	36: 2,	67: 2,	4: 2,)2: 2,	L0774: 2,	L0438: 2,	39: 2,
H0575: 3, H0036: 3,	S0010: 3, H0052:	S0438: 3, H0547: 3,	S0328: 3, S0380: 3,	S0206: 3, L0439: 3,	L0757: 3, L0595: 3,	H0423: 3, S0114: 2,	S0116: 2, S0212: 2,	H0484: 2, H0125: 2,	S0418: 2, S0376: 2,	S0444: 2, H0580: 2,	H0393: 2, H0369: 2,	H0370: 2, H0409: 2,	10331: 2, HO	H0486: 2, H0156: 2,	H0042: 2, H0318: 2,	H0581: 2, H0196: 2,	H0012: 2, L0163: 2,	H0083: 2, H0266: 2,	H0328: 2, H0615: 2,	J0194: 2, L0483: 2,	H0673: 2, H0169: 2,	S0364: 2, H0316: 2,	H0598: 2, S0036: 2,	H0090: 2, T0067: 2,	S0150: 2, S0344: 2,	S0422: 2, S0002: 2,	.0762: 2, L07	.0666: 2, L04	H0659: 2, H0539: 2,
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S0152: 2, H0521: 2,	S301	L0749: 2, L0756: 2,	S0434: 2, L0593: 2,	.0594: 2, L0366: 2,	10653: 2, H0543:	H0506: 2, H0171:	H0685: 1, H0716:	S0134	H0583	H0254: 1, H0255: 1	H0664:	10305: 1, L0005:	H0208:	S0046: 1	H0351:	10411: 1, H0261:	H0455:	H0632: 1	L0586:	10098: 1, H0120: 1,	S0182	00049: 1, H0230:	T011	H0545:	H0178:	H0123:	H0275:	H0239:	H0375: 1, H0288:
52: 2.	96: 2,	49: 2,	34: 2,	24: 2,	53: 2,	06: 2,	85: 1,	49: 1,	[8: 1,	54: 1,	10671: 1,	05: 1,	30468: 1,	50045: 1,	36026: 1,	11: 1,	10431: 1,	H0586: 1,	70039: 1,	98: 1,	48: 1,	<u>_</u>	H0596: 1,	H0546: 1,	H0009: 1,		H0024: 1,	r0010: 1,	75: 1,
S01;	90H	2	S04	105	H06	H05	H06	<u>100</u>	S02	H02	H06	H03	S046	800 _K	S602	H04	H04	H05	<u>1</u> 6	H00	100	S0049:	H05	H05	HOO	H0572:	H00.	<u>T</u> 00	H03
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H0286: 1, H0070:	H0622: 1, T0023: 1	S0368: 1, T0006:	H0553: 1, H0628:	H0032: 1, H0674:	H0124: 1, H0068:	H0038: 1, H0634:	H0412: 1, H0100:	560: 1, HC	561: 1, HC	H0130: 1, H0646: 1	S0142: 1, H0142:	H0695: 1, L0369:	.0371: 1, L0373:	.0764: 1, L0773:	0648: 1, LO	.0649: 1, L0784:	.0776: 1, L0558:	.0382: 1, L0787:	.0792: 1, LO	.0665: 1, H0520:	H0519: 1, H0593:	H0689: 1, H0683:	H0684: 1, H0435:	H0660: 1, H0648:	S0330: 1, H0710: 1	H0522: 1, S0044: 1	S0146: 1, S0406: 1,	H0555: 1, H0436:	H0447: 1, H0627: 1
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	<u> </u>	<u> </u>	<u> </u>		Pro-26 to Asn-34,	Arg-64 to Val-69	Val-97 to Lys-103.			Arg-10 to Ser-16.																		
		<u> </u>		_	3326			3327	3328	3329	,			_													3330	
	 			1 - 357	3 - 395			208 - 522	2 - 583	11 - 187			_														1 - 126	┙
			-	1150	1151			1152	1153	1154	-			-													1155	
				HCOOF77R	HCOQG37 HCOQG37R			HCOOG47R	HCOOH46R	TICOCITA HOOOTIAR	VITINOOU I													-			HCOO167R	-1
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		L0752: 4, H0156: 2,	L0803: 2, L0776: 2,	H0648: 2, L0439: 2,	H0431: 1, H0575: 1,	H0014: 1, T0023: 1,	H0598: 1, H0591: 1,	H0059: 1, L0772: 1,	L0372: 1, L0764: 1,	L0771: 1, L0773: 1,	L0804: 1, L0774: 1,	L0775: 1, L0375: 1,	L0805: 1, L0629: 1,	L0789: 1, L0665: 1,	L0438: 1, H0658: 1,	H0670: 1 and L0779: 1.									L0665: 4, L0743: 4,	H0486: 3, H0123: 3,	H0024: 3, L0528: 3,	L0748: 3, H0170: 2,
													,	٠			,			Arg-1 to Val-23.	Phe-94 to Glu-102,	Gln-133 to Thr-140.			Asn-29 to Gly-38,	Glu-50 to Gln-58.		
3331	3332	3333															3334	3335	3336	3337	3338		3339	3340	3341			
8 - 457	3 - 707	149 - 328		. :	:		_										1 - 471	1 - 519	3 - 398	3 - 143	2-613		3 - 149	75 - 443	2 - 430		-	
1156	1157	1158															1159	1160	1161	1162	1163		1164	1165	1166			
HCOQJ06R	HCOQJ07R	HCOQ136R										-					HCOQJ79 HCOQJ79R	HCOOK86R	HCOQL87R	HCOQM87R	HCOQ079 HCOQ079R 1		HCOQP32 HCOQP32R	HCOQS11R	HCOQUIS HCOQUISR			
НСООЛ06	HCOQJ07	HC0Q136															HCOQJ79	HCOQK86	HCOQL87	HCOQM87	НСООО79		HCOQP32	HCOQS11	HCOQU15			

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H0050: 2, L0769: 2,	H0670: 2, L0747: 2,	L0750: 2, H0171: 1,	H0556: 1, H0583: 1,	H0657: 1, S0212: 1,	H0208: 1, H0619: 1,	S0278: 1, H0550: 1,	S0222: 1, H0438: 1,	H0613: 1, H0331: 1,	H0013: 1, H0635: 1,	H0002: 1, S0474: 1,	H0581: 1, H0009: 1,	L0157: 1, H0012: 1,	S0025: 1, H0239: 1,	H0375: 1, H0266: 1,	H0598: 1, H0135: 1,	H0163: 1, H0087: 1,	H0551: 1, H0494: 1,	S0210: 1, L0640: 1,	L0763: 1, L0646: 1,	L0387: 1, L0766: 1,	L0805; 1, L0776; 1,	L0655: 1, L0657: 1,	H0144: 1, H0520: 1,	H0547: 1, H0519: 1,	H0689: 1, H0658: 1,	H0576: 1, L0779: 1 and	L0759: 1.	L0805: 7, L0776: 4, L0748: 4, H0545: 3,
				٠.		,							~							-								Ile-3 to Arg-8, Pro-17 to Ser-22,
																					,							3342
	•																											46 - 315
																									·	-		1167
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10641: 3,	.0592: 3,	10549: 2,	10486: 2,	10169: 2,	,0520: 2,	.0768: 2,	.0655: 2,	.0783: 2,	10144: 2,	10670: 2,	0406: 2,	0026: 2,	10170: 1,	0212: 1,	10125: 1,	(0580: 1,	10411: 1,	L0586: 1,	10390: 1,	, H0318: 1,	10620: 1,	,0006: 1,	10429: 1,	0386: 1,	0769: 1,	0374: 1,	L0662: 1,	0803: 1,	0526: 1,
S0438: 3, H0641: 3,	L0588: 3, L0592: 3,	T0049: 2, H0549: 2,	H0431: 2, H0486: 2,	L0471: 2, H0169: 2,	S0372: 2, L0520: 2,	L0553: 2, L0768: 2,	L0649: 2, L0655: 2,	L0527: 2, L0783: 2,	L0532: 2, H0144: 2,	H0682: 2, H0670: 2,	S0378: 2, S0406: 2,	1.0750: 2, S0026: 2,	H0542: 2, H0170: 1	H0716: 1, S0212:	H0638: 1, H0125: 1,	S0444: 1, H0580:	H0393: 1, H0411:	H0587: 1, 1	T0114: 1, H0390:	T0048: 1, H	H0596: 1, H0620:	H0375: 1, T0006: 1	H0606: 1, H0429:	S0294: 1, H0386: 1,	L0770: 1, L0769: 1	L3904: 1, L0374:	L0773: 1, L	L0766: 1, L0803:	L0774: 1, L0526:
Ser-28 to Ser-34,	Thr-45 to Thr-56,	Ser-58 to Thr-63,	Tyr-74 to Thr-79.																										
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L0809: 1, L0793: 1, L0664: 1, S0053: 1, H0689: 1, H0684: 1, H0659: 1, H0658: 1, H0648: 1, S0330: 1, H0521: 1, H0522: 1, S0404: 1, L0749: 1, L0780: 1, L0731: 1 and H0543: 1.	H0250: 35, L0770: 9, H0635: 6, L0483: 6, L0745: 6, R0521: 6, L0745: 6, S0003: 5, H0623: 5, L0776: 5, H0641: 4, L0598: 4, H0641: 4, L0598: 4, H0640: 3, H0656: 3, H0687: 3, H0669: 3, L0775: 3, L0665: 3, L0775: 3, L0665: 3, L0775: 3, L0665: 3, L0777: 3, S0134: 2, H0663: 2, S0007: 2, H0663: 2, H0613: 2, H0581: 2, H0613: 2, H0674: 2, S0440: 2, S0214: 2, H0613: 2, H0674: 2, S0440: 2,
	Glu-1 to Lys-17, Val-25 to Asn-30, Phe-36 to Pro-42.
	3343
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	1168
	HCOQV27R
	HC0QV27

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.0771: 2, L.0653: 2,	H0658: 2, H0710: 2,	10518: 2, S0206: 2,	L0754:	.0750: 2, L0779: 2,	.0731: 2, S0434: 2,	L0485:	J0362: 2, L0366: 2,	H0423	S0114:	F0049: 1, S0218: 1,	H0346:	H0661: 1, H0638: 1,	30356:	30358:	H0580:	H0645:	L0717: 1, H0431: 1,	H0609: 1, H0587: 1	H0427:	S0280: 1, L0021: 1,	H0004:	H0309: 1,	H0596: 1, H0024: 1	H0179: 1	H0271: 1, H0416: 1,	S0336: 1, S0316:	H0328: 1, H0622: 1	H0553: 1, H0032: 1	H0124: 1, H0708: 1
71: 2,	558: 2,	18: 2,	4:2,	50: 2,	31: 2,	05: 2,]	62: 2,]	43: 2,	70: 1,	49: 1,	02: 1,]	61: 1,	18: 1, 5	54: 1, 5	76: 1, 1	19: 1,	17: 1,]	09: 1,	H0632: 1,	S0280: 1, I	.69: 1,	S0049: 1, I	96: 1,	S0362: 1, I	71: 1,	36: 1, 5	28: 1,]	53: 1,]	24: 1,
L07	HOE	HO	102	<u>107</u>	107	200	<u> 1</u>	HOS	HOI	T00	9	HOH	S04	S03	803	H06	<u>1</u>	HOG	HOE	S02	HOS	800	H05	S03	H02	803	H03	HOS	띮
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S0366: 1. H0087: 1.	F0067: 1, H0100: 1,	0294: 1, H0633: 1,	H0649: 1, S0422: 1,	JNKWN: 1, L0667: 1,	.0641: 1, L0764: 1,	.0794: 1, L0649: 1,	.0774: 1, L0784: 1,	L0805: 1, L0657: 1,	.0659: 1, L0791: 1,	.0792: 1, L0664: 1,	S0052: 1, S0428: 1,	H0699: 1, H0703: 1,	H0693: 1, L0438: 1,	H0547: 1, H0593: 1,	H0659; 1, H0670: 1,	H0522: 1, S0013: 1,	H0555: 1, H0436: 1,	H0576: 1, H0478: 1,	S3014: 1, L0740: 1,	L0746: 1, L0758: 1,	L0588: 1, L0595: 1,	L0361: 1, S0026: 1,	S0242: 1, S0194: 1 and	H0352: 1.			S0282: 1, L0021: 1,	S0438: 1, L0775: 1,	L0666: 1, L0663: 1,
S		<u>.</u> <u>S</u>		رسر	<u>, p1</u>	_=		_==																	Trp-2 to Gly-7.				
					•																				3344	3345	3346		
																									23 - 439	1 - 321	271 510	016 - 1/7	-
			_							-															1169		111	1/11	
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H0659: 1, H0670: 1 and	LU/30. 1.				,		L0770: 3, H0685: 1,	H0581: 1, H0553: 1,	S0150: 1 and H0670: 1.	L0754: 14, H0052: 13,	L0748: 13, H0556: 12,	L0591: 12, L0755: 11,	L0752: 10, L0731: 9,	L0601: 9, L0439: 8,	L0777: 8, L0596: 8,	H0046: 7, H0521: 7,	L0758: 7, H0599: 6,	H0622: 6, S0002: 6,	L0646: 6, L0593: 6,	H0265: 5, S0474: 5,	L0776: 5, L0666: 5,	L0741: 5, L0747: 5,	S0026: 5, H0543: 5,	H0619: 4, S0222: 4,	H0013: 4, H0590: 4,	H0544: 4, H0620: 4,	H0124: 4, H0038: 4,
	Lys-1 to Lys-13.		3349. Pro-1 to Phe-15.	Val-1 to Gly-9.		Asn-1 to Gly-8.	Asn-7 to Pro-33,	Gly-40 to Gly-46,	Tyr-49 to Tyr-71.	Leu-35 to Tyr-40.	•						-							-		-	
	3347	3348	3349.	3350	3351	3352	3353			3354																	
	387 - 473	1 - 507	2 - 136	14 - 292	1 - 303	3 - 611	105 - 317			3 - 419														<u> </u>			
	1172	1173	1174	1175	1176	1177	1178			1179							-										
	HCOQZ02R	HCOQZ86 HCOQZ86R	HCORA95R	HCORB20R	HCORB43 HCORB43R	HCORB66R	HCORI18R			HCOR125R																	
	HCOQZ02	HC0QZ86	HCORA95	HCORB20	HCORB43	HCORB66	HCORI18			HCOR125													-				

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S0386: 4, LU/66: 4,	.0659: 4, H0547: 4,	10670: 4, S0378: 4,	.0740: 4, L0749: 4,	50: 4, S0434: 4,	.0588: 4, L0589: 4,	S0040: 3, H0580: 3,	S6026: 3, H0486: 3,	S0010: 3, S0051: 3,	10010: 3, H0083: 3,	H0266: 3, S0250: 3,	H0031: 3, H0591: 3,	S0142: 3, S0344: 3,	H0529: 3, L0770: 3,	.0769: 3, L0761: 3,	.0662: 3, L0774: 3,	.0653: 3, L0663: 3,	.0665: 3, L0438: 3,	H0435: 3, S0152: 3,	H0696: 3, S0342: 2,	F0049: 2, H0657: 2,	S0212: 2, H0638: 2,	S0418: 2, S0354: 2,	S0444: 2, S0360: 2,	S0046: 2, S0132: 2,	S0476: 2, L0717: 2,	S0278: 2, H0549: 2,	H0438: 2, H0587: 2,	H0333: 2, H0427: 2,	H0706: 2, H0545: 2,
850 <u>5</u>	1065	90H	L074	L075	1058	S004	S602	S001	T001	H026	H00:	S014	H05;) 1007) 1.066	<u>106</u>	1066	H04.	90H	T007	S021	S041	S04	800 <u>x</u>	S047	502	H04.	. H03.)LOH
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10012: 2	10014: 2,	10188:2	10553: 2	30364: 2,	10063: 2,	10059: 2	0042: 2,	10561: 2	10641:2	.0764: 2,	.0775: 2,	.0517: 2,	L0664: 2,	10519: 2,	10631:2	3014: 2,	.0757: 2,	0599: 2,	.0362: 2,	10653: 2,	10667:2	10423: 2	70002: 1,	10686: 1	10255: 1	10663: 1	10458: 1	0442: 1,	0208: 1,
H0050: 2, H0012: 2,	S0050: 2, H0014: 2,	S6028: 2, H0188: 2,	H0604: 2, H0553: 2,	H0617: 2, S0364: 2,	S0366: 2, H0063: 2,	H0087: 2, H0059: 2,	S0038: 2, T0042: 2,	H0560: 2, H0561: 2,	H0509: 2, H0641: 2,	H0647: 2, L0764: 2,	.0771: 2, L0775: 2,	.0655: 2, L0517: 2,	.0783: 2, L	.0565: 2, H0519: 2,	H0658: 2, H0631: 2,	S0390: 2, S3014: 2,	J0745: 2, L0757: 2,	S0436: 2, L0599: 2,	0595: 2, L0362: 2,	S0011: 2, H0653: 2	H0665: 2, H0667: 2,	H0542: 2, H0423: 2,	H0422: 2, T0002: 1	H0222: 1, H0686:	H0717: 1, H0255:	H0661: 1, H0663:	H0662: 1, H0458:	S0420: 1, S0442:	S0376: 1, H0208: 1
H	80	9S	Ħ	Œ	<u>8</u>	Ħ	<u>S</u>	Ħ	虽	Ħ	크	<u> </u>	<u> </u>	<u> </u>	H	<u>S</u>	ន	20	<u>3</u>	SO	H	H	H	呈	H	H	H	80	80
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S0045: 1, H0645: 1	S0300: 1, H0406: 1,	S0220: 1, H043	H0455: 1, H040	H0600: 1, H058	H0642: 1, H049	T0109: 1, H0069: 1,	S0280: 1, H015	H0004: 1, H06	H0253: 1, S004	H0194: 1, H02.	T0103: 1, T0115: 1	H0327: 1, H0457: 1	L0041: 1, H056	L0471: 1, H01:	H0015: 1, S0362: 1	H0239: 1, H0355: 1	H0510: 1, H03	H0594: 1, S0022: 1	H0428: 1, H06	H0628: 1, H00	L0456: 1, S003	H0040: 1, H0616: 1,	H0551:-1, H02	H0056: 1, H06	H0100: 1, H0494: 1,	S0440: 1, H0131:	H0130: 1, H0633:	S0422: 1, S0426:	L0369: 1, L3905: 1
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L0667: 1, L0363: 1,	L0794: 1, L0803: 1,	L0375: 1, L0651: 1,	L0806: 1, L0805: 1,	L0654: 1, L0379: 1,	L0526: 1, L0519: 1,	L0647: 1, L0791: 1,	S0428: 1, H0144: 1,	S0374: 1, H0520: 1,	S0126: 1, H0711: 1,	H0660: 1, H0672: 1,	S0328: 1, H0539: 1,	S0406: 1, H0555: 1,	H0436: 1, H0627: 1,	S0432: 1, S0037: 1,	L0743: 1, L0744: 1,	L0751: 1, S0031: 1,	S0394: 1, L0592: 1,	L0485: 1, L0608: 1,	L0366: 1, H0136: 1,	H0506: 1 and H0352:			•						
																					Glu-9 to Gly-17.	Lys-14 to Gly-26,	Glu-33 to Arg-44,	Pro-64 to Pro-69,	Pro-79 to Asn-88,	Thr-109 to Pro-123.		Ala-17 to Asn-23.	Ser-1 to Thr-8,
											•										3355	3356					3357	3358	3359
																					48 - 206	1 - 537					2 - 463	158 - 325	37 - 303
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							•														HCQCR82R	HCRME42R				•	HDABR53R	HDLBE03R	HDPF126R
																					HCQCR82	HCRME42					HDABR53	HDLBE03	HDPF126

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:	L0759: 10, L0774: 3, H0648: 3, H0686: 2,	S0360: 2, L0622: 2,	H0486: 2, L0521: 2,	L0775: 2, H0659: 2,	S0328: 2, L0779: 2,	L0755: 2, H0341: 1,	S0354: 1, S0358: 1,	H0351: 1, S0222: 1,	L0021: 1, H0025: 1,	L0105: 1, T0115: 1,	H0545: 1, H0571: 1,	H0050: 1, L0471: 1,	S6028: 1, S0003: 1,	H0031: 1, H0316: 1,	H0038: 1, H0087: 1,	H0494: 1, H0509: 1,	S0142: 1, L0520: 1,	L0769: 1, L0638: 1,	L0772: 1, L0794: 1,	L0803: 1, L0804: 1,	L0382: 1, L0367: 1,	L0665: 1, S0374: 1,	H0690: 1, H0684: 1,	S0044: 1, S3012: 1,	S0027: 1, L0744: 1,	L0777: 1, L0758: 1,	L0604: 1, S0026: 1 and
Glu-17 to Ala-32, Arg-39 to Trp-47.			٠																							-	
	3360		•																								
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S0276: 1.					L0748: 50, L0754: 42,	L0731: 31, H0644: 30,	L0747: 29, H0551: 22,	S0418: 21, L0659: 17,	H0619: 15, L0663: 14,	L0666: 13, S0360: 10,	H0622: 10, H0031: 10,	H0553: 10, L0662: 10,	S0126: 10, L0757: 10,	L0603: 10, S0420: 8,	H0486: 8, H0013: 8,	H0050: 8, H0623: 8,	L0717: 7, H0144: 7,	L0770: 6, L0750: 6,	H0427: 5, H0024: 5,	H0328: 5, H0030: 5,	L0773: 5, L0665: 5,	L0749: 5, H0713: 4,	T0039: 4, S0280: 4,	H0012: 4, H0038: 4,	L0794: 4, L0774: 4,	L0775: 4, L0805: 4,	H0624: 3, H0170: 3,	H0171: 3, S0356: 3,	H0329: 3, S0046: 3,
	Leu-21 to Ala-27,	Pro-48 to Gly-55.	Ala-14 to Leu-20.	Tyr-25 to Phe-32.	-																								
	3361		3362	3363	3364																								
	231 - 431		207 - 431	102 - 215	179 - 364					-								_											
	1186	00.	118/	1188	1189						*****															 -		-	
	HDTBP08R		HDIDAZIK	HDTDB88R	HDTJR38R														-										·
	HDTBP08	1100000	HDI DAZI	HDTDB88	HDTJR38								_																

110221.3 H0575.3	0471: 3, HOS70: 3,	504/1: 3, 110025: 3, 50250: 3 H0032: 3.	H0591: 3, L0598: 3,	0803: 3, L0653: 3,	.0657: 3, H0670: 3,	S0406: 3, H0555: 3,	.0780: 3, L0605: 3,	H0667: 3, H0717: 2,	F0049: 2, S0212: 2,	H0661: 2, S0376: 2,	S0007: 2, H0208: 2,	S0476: 2, H0645: 2,	H0411: 2, H0437: 2,	10550: 2, H0586: 2,	H0333; 2, H0599; 2,	H0042: 2, H0545: 2,	H0123: 2, H0105: 2,	H0014: 2, H0051: 2,	H0375: 2, L0142: 2,	H0598: 2, H0135: 2,		: 2, L0640: 2,	: 2, L5566: 2,	: 2, L0646: 2,	: 2, L0363: 2,	0768: 2, L0375: 2,	.; 2, L0776: 2,	0809: 2, L0438: 2,	7: 2, H0519: 2,
1102210	1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1		H0591:	L0803:	L0657:	S0406:	L0780:	H0667:	T0049:	H0661:	30007:	S0476:	H0411:	H0550:	H0333:	H0042:	H0123:	H0014:	H0375	H0598	H0040	1.0520: 2,	L0769: 2,	1.0772: 2,	L0771: 2,	T0768	L0654: 2, 1	60807	H0547: 2,
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L0602: 2, L0743: 2,	L0744: 2, L0439: 2,	L0758: 2, L0759: 2,	L0588: 2, L0599: 2,	L0361: 2, H0716: 1,	H0294: 1, S0430: 1,	S0298: 1, S0132: 1,	H0351: 1, H0369: 1,	S0222: 1, H0370: 1,	H0392: 1, H0409: 1,	H0587: 1, H0574: 1,	H0485: 1, T0040: 1,	•	L0021: 1, H0003: 1,	H0036: 1, H0274: 1,	H0253: 1, H0505: 1,	H0251: 1, H0544: 1,	H0546: 1, H0046: 1,	H0081: 1, H0019: 1,	F0003: 1, H0057: 1,	H0015: 1, H0373: 1,	H0275: 1, H0594: 1,	S0003: 1, H0039: 1,	H0111: 1, H0674: 1,	S0364: 1, S0036: 1,	H0090: 1, H0063: 1,	H0087: 1, H0268: 1,	H0412: 1, H0059: 1,	H0100: 1, L0564: 1,	H0561: 1, S0210: 1,
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L0369: 1, L3904: 1, L0800: 1, L0641: 1, L0764: 1, L5564: 1, L0804: 1, L0378: 1, L0806: 1, L0807: 1, L0526: 1, L0787: 1, L0788: 1, L0791: 1, L0793: 1, S0374: 1, H0691: 1, T0068: 1, H0690: 1, H0682: 1, H0690: 1, H0698: 1, H0672: 1, H0651: 1, S0146: 1, S0037: 1, S0146: 1, S0027: 1, S0032: 1, H0445: 1, L0589: 1, L0590: 1, L0589: 1, L0590: 1, L0485: 1, L0604: 1, H0668: 1, H0653: 1, S0242: 1, S0194: 1 and H0008: 1,		·		
	Thr-8 to Thr-22, Phe-42 to Lys-51, Arg-60 to Lys-69.	Arg-1 to Cys-23, Trp-25 to Ser-35.	Ser-10 to Thr-34, Ser-37 to Asn-89.	Thr-15 to Ser-26.
	3365	3366	3367	3368
	1 - 288	3 - 170	32 - 376	3 - 128
	1190	1191	1192	1193
	HE8QX44R	HE9QU94R	HEAHF02R	HEEAY40R
	HE8QX44	HE9QU94	HEAHF02	HEEAY40

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	H0457; 14, L0751: 13,	H0617: 8, L0761: 7,	L0666: 6, L0664: 6,	L0665: 6, H0593: 6,	L0662: 5, L0748: 5,	H0052: 4, H0014: 4,	H0529: 4, L0655: 4,	L0809: 4, L0663: 4,	H0436: 4, L0747: 4,	L0758: 4, H0592: 3,	H0587: 3, H0581: 3,	L0763: 3, L0769: 3,	L0659: 3, H0539: 3,	L0779: 3, S0434: 3,	H0506: 3, H0295: 2,	H0650: 2, H0657: 2,	L0005: 2, S0356: 2,	S0442: 2, S0354: 2,	S0360: 2, H0580: 2,	H0733: 2, H0553: 2,	H0135: 2, H0560: 2,	H0625: 2, H0647: 2,	L0374: 2, L0657: 2,	L0647: 2, H0682: 2,	H0672: 2, H0521: 2,	S0028: 2, L0743: 2,	L0749: 2, L0752: 2,	L0755: 2, L0759: 2,	L0593: 2, S0446: 2,
Glu-9 to Gly-17.	Pro-13 to Leu-25.			-			-														-			•					
3369	3370																												
45 - 221	96 - 395																			-					•				
1194	1195			•	'					*******						-								-					
HEGAF68R	HEOSP11R									-																			
HEGAF68	HEOSP11																												

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.0600: 2, H0556: 1	7:1	1:1	33: 1	8: 1	S0420: 1, S0358: 1,	 	S0045: 1, S0132: 1,	1.6	1:1	00.1	93.	3: 1	S0280: 1, H0599: 1,	8: 1	H0318: 1, S0474: 1,	1:1	5: 1	57: 1	0.5	, S0051: 1,	7:1	2.7	3:1	1:1	6: 1	4: 1	2: 1	 i	0:1,
1055	H0686: 1, H0717: 1	000	106	H0402: 1, S0418: 1	035	8	013	1054	<u>₹</u>	1060	1055	1001	059	1061	740	1025	524	1056	100£	905	890	H0622:	[021	[018	, H0056: 1	545	H0561: 1, H0652: 1	934	.0625: 1, L0770: 1
2, F	1, F	1,5	1, I	1,5	1, S	1, S	1, S	1, F	1, H	1,1	1, F	1	Œ	. H	 	1, H	1, H	1,1	1, F	1, S	1, H	1,1	1, H	1, H	1,1	1, E	1, H	1, S(1, L
8	586:	556:	561:	1 02;	30	76:	45:	519:	22:	392:	386:	186:	80:	575:	318:	6	15:	99	571:	,0471: 1,	36028: 1,	H0428: 1,	70006: 1,	.0142: 1,	H0032: 1,	H0100: 1,	61;	45	25:
ន្ន	Ĕ	<u>Ř</u>	<u>ĕ</u>	Ě	8	<u>8</u>	<u>8</u>	<u>H</u>	S02	H	HOH	<u>H</u>	S02	HOH	豆	200	<u>10</u>	<u>H</u>	<u>10</u>	7	860	<u> 1</u>	<u>1</u> 00	<u> </u>	<u>H</u>	<u>E</u>	H05	S01	9
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L0639: 1, L0646: 1, L0800: 1, L0641: 1, L0642: 1, L0648: 1, L0794: 1, L0766: 1, L0679: 1, L0766: 1, L0629: 1, L0651: 1, L0629: 1, L0651: 1, L0629: 1, L0656: 1, L0792: 1, S0428: 1, H0726: 1, S0428: 1, H0690: 1, H0684: 1, H0658: 1, L0611: 1, S0404: 1, S0406: 1, H0555: 1, L0771: 1, S0436: 1, L0777: 1, S0436: 1, L0601: 1, L0366: 1, S0106: 1,					H0594: 18, L0777: 10,	L0766: 6, H0595: 6, H0170: 5, H0373: 5,
	Asn-4 to Tyr-9.	Ala-16 to Arg-26, Pro-68 to Gly-78.				
	3371	3372	3373	3374	3375	
	26 - 112	102 - 365	2 - 409	32 - 151	2 - 211	-
	1196	1197	1198	1199	1200	
	HFABK01R	HFADS75R	HFIBG63R	HFIJB15R	HFIXK57R	
	HFABK01	HFADS75	HFIBG63	HFIJB15	HFIXK57	

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)]) <u>S</u>	<u>ਹ</u>	<u> </u>	H	SC	<u> </u>	<u>구</u>	<u> </u>		S				<u>×</u>	<u> </u>	<u> </u>	S	<u> </u>	<u> </u>	<u> </u>	<u> </u>	프	<u> </u>	<u> </u>		<u> </u>	<u> </u>	<u> </u>	14
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L0761: 1, L0667: 1,	L0764: 1, L0521: 1,	L0533: 1, L0775: 1,	L0651: 1, L0806: 1,	L0655: 1, L0656: 1,	S0374: 1, H0547: 1,	S0328: 1, H0539: 1,	H0696: 1, L0741: 1,	L0744: 1, L0740: 1,	L0754: 1, L0747: 1,	L0750: 1, L0753: 1,	L0759: 1, S0031: 1,	L0485: 1, L0362: 1,	S0242: 1, H0543: 1 and	S0398: 1.				AR089: 4, AR096: 3,	AR033: 3, AR060: 3,		AR055: 1, AR053: 0,	0, AR052:	S0222: 1, S0049: 1,	H0272: 1, H0670: 1,	H0521: 1, H0555: 1 and	S0436: 1.	
															Glu-27 to Leu-36.	Pro-3 to Tyr-22.	Lys-17 to Thr-23.	Pro-75 to Cys-89.	•						-		Cys-5 to Pro-21,
															3376	3377	3378	3379			-			•			3380
			•						•						18 - 137	3 - 86	75 - 233	1 - 270									18 - 314
				-											1201	1202	1203	1204					 	-			1205
			-												HFIZQ64R	HFKKK36R		HFPIX37R		-							HFTDK36 HFTDK36R
															HFIZQ64	HFKKK36	HFPEC93	HFPIX37									HFTDK36

							•																						
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His-66 to Leu-77,	Gln-80 to Ala-88,	Tyr-92 to Ser-98.			Val-24 to Leu-31.			Arg-2 to Ser-7.																					
			3381	3382	3383	3384	3385	3386	3387																				
			11 - 250	353 - 57	304 - 426	54 - 230	1 - 474	53 - 277	181 - 426																				
			1206	1207	1208	1209	1210	1211	1212	1											<u></u>								
			HFVIB28R	HFXGR60R	HHAUD07R	HHBFO21 HHBFO21R	HHEVG50R	HHFGO65 HHFGO65RA	ниеов48 ингов48кр	00B	1							-											
			HFVIB28	HFXGR60	HHAUD07	HHBF021	HHEVG50	HHFG065	HHFOE48	! ! !																			

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H0252: 2, H0428: 2	H0628: 2, H0551: 2,	S0210: 2, L0763: 2,	.0770: 2, L0774: 2,	2, L0518: 2,	.0809: 2, H0547: 2,	H0682: 2, H0670: 2,	S0037: 2, S0027: 2,	.0751: 2, L0752: 2,	.0758: 2, L0601: 2,	H0668: 2, S0194: 2,	H0170: 1, H0556: 1	H0686: 1, S0040: 1,	H0716: 1, H0295: 1,	H0341: 1, H0638: 1,	S0418: 1, S0376: 1,	, H0580: 1,	H0329: 1, S0468: 1,	, S0046: 1,	, S0476: 1,	H0619: 1, H0645: 1,	, H0549: 1,	1, H0586: 1,	l, L0021: 1,	1, H0581: 1,	H0052: 1, H0309: 1,	H0546: 1, H0457: 1,	H0150: 1, H0041: 1,	H0563: 1, H0050: 1,	.0163: 1, H0051: 1,
H0252:	H0628:	S0210: 3	L0770: 3	L0661: 2,	L0809:	H0682:	S0037: 2	L0751: 3	L0758: 2	H0668:	H0170:	H0686:	H0716:	H0341:	S0418: 1	S0444: 1,	H0329:	S0045: 1,	S0132: 1,	H0619:	L0717: 1,	H0550: 1,	H0587: 1,	H0575: 1,	H0052: 1	H0546: 1	H0150:]	H0563:	L0163: 1
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H0615: 1, H0688: 1,	H0031: 1, H0634: 1,	H0087: 1, H0334: 1,	H0633: 1, H0646: 1,	L0772: 1, L0800: 1,	L0643: 1, L0764: 1,	L0662: 1, L0767: 1,	L0775: 1, L0651: 1,	L0806: 1, L0805: 1,	L0776: 1, L0656: 1,	L0783: 1, L0383: 1,	L0543: 1, L0789: 1,	L0663: 1, H0593: 1,	₩	_	H0709: 1, S0152: 1,	H0521: 1, H0627: 1,	L0611: 1, L0439: 1,	L0745: 1, L0759: 1,	L0593: 1, L0361: 1,	L0603: 1, S0026: 1,	H0667: 1 and H0506: 1.								L0752: 11, L0662: 9,
				•																					Pro-13 to Leu-18,	Leu-37 to Arg-42,	Lys-65 to Gly-77.		Val-40 to Tyr-46.
	_														-							3388	3389	3390	3391			3392	3393
			•			-											-					3 - 62	77 - 142	112 - 270	49 - 315			244 - 405	1 - 138
				_																		1213	1214	1215	1216			1217	1218
															,							HHSFG15R	HHSGP15R	HHSG017R	HKBAD05 HKBAD05R	-		HKZAD21 HKZAD21R	HKZAE07R
																	•					HHSFG15	HHSGP15	HHSG017	HKBAD05			HKZAD21	HKZAE07

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.0596: 9, L0659: 8,	.0665: 7, L0766: 5,	.0750: 5, L0521: 4,	.0740: 4, L.0757: 4,	0771: 3, L0775: 3,	.0809; 3, L0666; 3,	.0664: 3, H0648: 3,	.0755: 3, L0758: 3,	.0591: 3, L0785: 2,	.0717: 2, H0036: 2,	H0052: 2, H0529: 2,	.0769: 2, L0637: 2,	.0646: 2, L0765: 2,	.0651: 2, L0653: 2,	.0776: 2, L0655: 2,	.0663: 2, H0682: 2,	.0748: 2, L0439: 2,	.0731: 2, L0759: 2,	.0593: 2, L0595: 2,	.0601: 2, H0686: 1,	S0134: 1, L0778: 1,	H0484: 1, H0661: 1,	H0662: 1, H0393: 1,	H0643: 1, H0002: 1,	H0599: 1, H0318: 1,	S0049: 1, H0251: 1,	_0040: 1, H0545: 1,	.0041: 1, N0006: 1,	N0007: 1, S0388: 1,	L0483: 1, H0606: 1,
F029	990T	F075	1.074	1.077	D80T	1066 1066	1.075	1059	1701	00H	9/07	1064	5907	101	9907	L074	1.073	F020	0907	5013	H048)90H	790H	H059	8004	1007	1007	000 <u>N</u>	L048
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H0038: 1, H0040: 1, H0551: 1, H0413: 1, H0538: 1, S0002: 1, L0763: 1, L0772: 1, L0364: 1, L0649: 1, L0542: 1, L0657: 1, L0542: 1, L0526: 1, L0783: 1, L0528: 1, L0788: 1, L0528: 1, H0689: 1, H0690: 1, H0689: 1, H0690: 1, H0660: 1, S0004: 1, S0044: 1, S0404: 1, L0751: 1, L0742: 1, L0751: 1, L0742: 1, L0751: 1, L0742: 1, L0751: 1, L0603: 1, H0543: 1, H0422: 1 and H0543: 1, H0422: 1 and H0543: 1, H0422: 1 and		
	Gly-1 to Val-7, Lys-15 to Leu-25, Pro-52 to Cys-57.	Lys-47 to Glu-52, Ala-71 to Asp-76, Asp-89 to Ser-110.
	3394	3395
	79 - 327	3 - 368
	1219	1220
	HKZAG45 HKZAG45RA	HKZA114R
	HKZAG45	HKZAI14

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S0474: 22, S0360: 13,	110556, 11 00250, 11	T 0747 11 H0087 10	50176: 10, 110134: 0	S0126: 10, H0124: 9,	S0410: 8, S0278: 8,	H0024: 8, L0588: 8,	\$0206: 7, \$0046: 6,	L0731: 6, H0543: 6,	H0265: 5, S0212: 5,	S0045: 5, H0620: 5,	H0594: 5, H0090: 5,	H0063: 5, H0551: 5,	T0042: 5, L0530: 5,	H0144: 5, H0696: 5,	S3014: 5, L0758: 5,	S0192: 5, S0116: 4,	H0549: 4, H0013: 4,	H0530: 4, H0544: 4,	H0081: 4, L0471: 4,	H0012: 4, H0644: 4,	H0163: 4, H0412: 4,	H0100: 4, L0769: 4,	L0771: 4, S0027: 4,	L0740: 4, L0757: 4,	L0605: 4, H0423: 4,	H0624: 3, H0713: 3,	S0418: 3, S0444: 3,	S0408: 3, H0251: 3,	H0545: 3, S0051: 3,	H0083: 3, H0290: 3,
3396 Ser-9 to Ala-15.	_																													
3396																				_										
2 - 79																				•										
1221																														
HKZAI68 HKZAI68RA																										,	•			
HKZAI68																														

F0041.3 S0344: 3.	0662: 3, L0768: 3,	,0806: 3, L0659: 3,	.0608: 3, S0194: 3,	H0352: 3, S0040: 2,	H0717: 2, H0716: 2,	S0134: 2, H0341: 2,	H0402: 2, H0305: 2,	S0356: 2, H0580: 2,	S0476; 2, H0411; 2,	H0392: 2, H0586: 2,	H0575; 2, H0052; 2,	H0327: 2, H0546: 2,	H0373: 2, L0163: 2,	H0687: 2, H0284: 2,	H0286: 2, H0428: 2,	H0039: 2, H0628: 2,	H0413: 2, H0056: 2,	S0440: 2, L0598: 2,	.0770: 2, L0521: 2,	776: 2, L0542: 2,	L0783: 2, L0529: 2,	0663: 2, H0519: 2,	10659: 2, S0378: 2,	H0518: 2, S0044: 2,	H0555: 2, H0627: 2,	S0028: 2, L0748: 2,	.0756: 2, L0755: 2,	S0031: 2, H0445: 2,	.0596: 2, L0591: 2,
TOUT	907	1080	106	H03	10H	501	H04	503	804	H03	HOS	HO3	HO	HOC	HO	HO	70H	70S			io.	07	OH	HO	OH HO	OS	<u> </u>	os	07
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S0026: 2	H0171:	S0342: 1	H0656: 1	H0484: 1	H0663:	SQ376: 1	S6022: 1	H0370:	H0333:	L0623: 1	T0114: 1	H0427:	H0156: 1	H0706: 1	H0581:	H0196: 1	H0235:	H0046: 1	H0041:	H0015:	N0007:	H0375:]	H0271:]	H0292: 1	H0622: 1	H0553: 1	H0032: 1	H0674: 1	H0616: 1
J0361: 2, S0026: 2,	10506: 2, H0171: 1	H0157: 1, S0342: 1	0049: 1, H0656: 1	0282: 1,	10255: 1, H0663: 1	S0420: 1, S0376: 1,	L0717: 1, S6022: 1,	0441: 1,	10592: 1, H0333: 1	H0492: 1, L0623:	70039: 1,	H0635: 1, H0427:	30280: 1, H0156:	.0021: 1, H0706:	H0004: 1, H0581:	S0049: 1,	10085: 1, H0235: 1,	r0115: 1, H0046: 1	H0086: 1, H0041: 1	H0123: 1, H0015:	10051: 1, N0007: 1	10408: 1, H0375:	H0266: 1,	H0416: 1, H0292:	S0003: 1, H0622:	.0483: 1,	H0617: 1,	H0673: 1,	H0135: 1, H0616:
1	正	<u> </u>	<u> </u>	S	<u> </u>	<u> </u>	<u> </u>	<u>æ</u>	<u> </u>	프	F	<u> </u>	<u>S</u>	7	Œ	<u></u> <u> </u>	<u> </u>	F	<u>E</u>	正	王	<u> </u>	王	<u> </u>	<u> </u>	크	<u>H</u>	<u> </u>	H
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)438: 1,	150: 1,)144: 1,	S0002: 1,	695: 1,	772: 1,	764: 1,	L0650: 1,	775: 1,	l, L0657: 1,	L0526: 1,	L0782: 1,	L0519: 1,	.0793: 1,	0702: 1,	0547: 1,	0682: 1,	0672: 1,	l, S0328: 1,	0539: 1,	0521: 1,	1, S0004: 1,	0436: 1,	390: 1,	, S0032: 1,	l, L0439: 1,	L0750: 1,	3759: 1,	0595: 1,	0604: 1,
H0623: 1, S0438: 1	H0132: 1, S0150: 1	H0633: 1, S0144: 1	S0210: 1, S0	S0426: 1, H0695: 1	L0763: 1, L0772: 1	L0643: 1, LC	L0766: 1, L0650: 1	L0774: 1, LC	L0654: 1, L(_	L0518: 1, L(L0666: 1, H	H0701: 1, H0547:	H0689: 1, H	H0660: 1, H0672:	H0651: 1, S(S0330: 1, H	S0380: 1, H	H0522: 1, SI	H0134: 1, H0436: 1	S0392: 1, S0390: 1	S0037: 1, SC	L0743: 1, L0	L0754: 1, L0	L0779: 1, L0759: 1	S0260: 1, H(H0707: 1, L0604: 1
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	Ser-1 to Lys-6, Asp-48 to Gly-55, Pro-68 to Phe-76, Pro-83 to Gly-89, Asn-100 to Asp- 108, Glu-134 to Lys-141, Lys-144 to Asp-160.	Met-41 to Gln-47.	Gln-54 to Leu-59, Glu-70 to Lys-76, Lys-147 to Ser-154.	Asp-1 to His-35, Pro-67 to His-75.	Ser-11 to Gly-18.
	3397	3398	3399	3400	3401
	1 - 480	3 - 266	1 - 462	1 - 276	2 - 103
	1222	1223	1224	1225	1226
		HKZAR58R	HKZAŚ59R	HKZAS64R	HKZAS84R
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1 C	, L	4, 1.07	3, T00	3, 1.06	3, H02	3, 101	3, HO	3, HO	3, 1.03	3, 1.07	3. 123	3, HO	3, 1.07	J0752: 3, S0260: 3,	3, 10	.0588: 3, L0599: 3,	2, H0	2, 50	2, S0	2, H0	H0392: 2, H0599: 2,	H0575:'2, H0052: 2,	H0309: 2, H0085: 2,	H0597: 2, H0544: 2,	H0428: 2, H0124: 2,	S0036: 2, H0163: 2,	H0038: 2, H0413: 2,	H0529: 2, L0369: 2,	.0763: 2, L0770: 2,	0638: 2, L0764: 2,
1	H0658: 4, L0/4/: 4	.0750: 4, L0757: 4,	H0295: 3, T0049: 3,	H0255: 3, L0622: 3	Γ0039: 3, H0253: 3	10545: 3, L0163:	H0594: 3, H0188: 3,	H0673: 3, H0100: 3,	S0142: 3, L0372: 3,	.0775: 3, L0776: 3,	0657: 3, L0382: 3,	,0665: 3, H0672: 3,	S0027: 3, L0742: 3,	.0752:	H0445: 3, L0596: 3,	.0588:	S0420: 2, H0637: 2,	S0045: 2, S0132: 2,	S0300: 2, S0278: 2,	S0222: 2, H0431: 2,	H0392	H0575	H0305	H0597	H0428	S0036	H003	H0529	1.076	1,063
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83	375	384	532	126	651	82	485	601	424	556	89	342	114	650	785	180	484	254	9662	638	356:	S0358;	208	438	905	L0477: 1,	S0280: 1	H0590: 1,	618
J0771: 2, L0803: 2	.0804: 2, L0375: 2,	.0655: 2, L0384: 2,	J0809: 2, L0532: 2,	80374: 2, \$0126: 2,	H0670: 2, H0651: 2,	H0521: 2, S0028: 2,	.0740: 2, L0485: 2,	J0362: 2, L0601: 2,	S0192: 2, S0424: 2,	JO600: 2, H0556: 1,	r0002: 1, H0686: 1,	30040: 1, S0342:	H0294: 1, S0114:	S0134: 1, H0650:	H0657: 1, L0785:	H0341: 1, S0180:	S0212: 1, H0484:	H0483: 1, H0254: 1	H0671: 1, H0662: 1	H0402: 1, H0638: 1	S0418: 1, S0356: 1	8	S0007: 1, H0208: 1	H0441: 1, H0438: 1	H0587: 1, S0005: 1,	ន្ទ	SS	出	H0004: 1, H0618: 1
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077	080	90	080	337	190	052	974	036	219	9	900	ğ	029	213	965	034	217	848	067	8	413	Γ0007: 1,	8	4	058	.0623: 1,	P0114: 1,	.0021: 1,	8
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١.	S0049: 1, H0251: 1 S0049: 1, H0251: 1	H0596: 1, T0110: 1,	H0327: 1, H000	H0123: 1, L047	H0006: 1, H001	H0051: 1, S038	H0510: 1, S033.	H0687: 1, T002	T0006: 1, H042	H0182: 1, H060	H0383: 1, H0165: 1	H0166: 1, S036	H0135: 1, H0090: 1	T0067: 1, H048	H0412: 1, T0069:	S0112: 1, L056	H0494: 1, S0015: 1,	H0561: 1, L076	L0640: 1, L063	L0761: 1, L0667: 1	L0772: 1, L0373: 1	L0646: 1, L064	•	L0773: 1, L0648: 1	L0521: 1, L0768:	L0364: 1, L053	L0650: 1, L0774: 1	L0376: 1, L0806: 1	L0652: 1, L0653:
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L0606: 1, L0629: 1, L0659: 1, L0636: 1, L0542: 1, L0636: 1, L0518: 1, L0782: 1, L0783: 1, H0689: 1, H0690: 1, H0689: 1, H0660: 1, S0328: 1, S0330: 1, S0328: 1, H0522: 1, S0004: 1, H0555: 1, L0611: 1, S3014: 1, S0206: 1, L0755: 1, L0758: 1, L0755: 1, L0608: 1, L0589: 1, L0608: 1, L0591: 1, L0608: 1, L0561: 1, L0366: 1,	H0423: 1, S0460: 1, S0456: 1 and H0506: 1.			L0771: 3, L0021: 1, S0036: 1, H0689: 1 and
		Val-6 to Lys-13, Tyr-23 to Asn-33, Lys-46 to Lys-51, Leu-75 to Asn-85, Asp-101 to Glu-106.	Cys-1 to Val-33, Ala-47 to Glu-54.	
,		3402	3403	3404
		2 - 412	2 - 163	137 - 361
		1227	1228	1229
		HKZAV69R	HKZAV72R	HKZAX48 HKZAX48R
·		HKZAV69	HKZAV72	HKZAX48

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H0659: 1.																			L0766: 5, L0751: 4,	H0662: 2, H0253: 2,	L0754: 2, L0750: 2,	H0170: 1, H0587: 1,	H0327: 1, H0688: 1,
	Arg-5 to Leu-11, Arg-32 to Lys-39, Thr-117 to Lys-122.	Gin.41 to His.48, Leu.55 to Gly.61, Pro.71 to Gly.79, Ala.104 to His-111.		Ala-2 to Lys-12.			Pro-77 to Lys-85.	Pro-82 to Lys-92.		Gin-20 to Ser-30.	Phe-45 to Ser-51.			Lys-17 to Thr-23.	Met-40 to Thr-52.		Lys-18 to Lys-30.	Lys-18 to Lys-30.	Pro-37 to Arg-46,	Gln-75 to Gly-81,	Pro-83 to Ser-89.		
	3405	3406	3407	3408	3409	3410	3411	3412	3413	3414	3415	3416	3417	3418	3419	3420	3421	3422	3423				
	2 - 520	1 - 381	2-112	2 - 190	2 - 151	430 - 648	144 - 398	118 - 411	331 - 585	2 - 337	57 - 275	2 - 238	131 - 199	67 - 225	3 - 161	324 - 145	136 - 276	2 - 196	127 - 474				
	1230	1231	1232	1233	1234	1235	1236	1237	1238	1239	1240	1241	1242	1243	1244	1245	1246	1247	1248				
	HKZBB22R	HKZBS89R	HLDQQ80R	HLDXE19R	HLICD55R	HLJB132R	HLJB137R	HLTHA47R	HLTJA50R	HLWAC95 HLWAC95R	HLWDZ38R	HLYDIS7 HLYDIS7R	HMCF019R	HMCIZ44R	HMCJE25R	HMSPB25 HMSPB25R	HMVBB04 HMVBB04R	HNAAE01 HNAAE01R	HNBUP08R				
	HKZBB22	HKZBS89	HLDQQ80	HLDXE19	HLICD55	HLJB132	HLJB137	HLTHA47	1	HLWAC95	HLWDZ38	HLYDI57	HMCF019	HMCIZ44		HMSPB25	HMVBB04	HNAAE01	HNBUP08				

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H0428: 1, S0210: 1, L0763: 1, L0770: 1, L0769: 1, L0646: 1, L0652: 1, L0803: 1, L0776: 1, L0659: 1, L0789: 1, L0666: 1, L0663: 1, L0665: 1, H0660: 1, S0380: 1, L0740: 1, L0755: 1, L0758: 1, L0589: 1, L0758: 1, L0589: 1,	L0783: 4, \$0280: 2, H0662: 1, H0550: 1, L0764: 1, L0662: 1, L0809: 1 and H0684: 1.						
H0428: 1, L0763: 1, L0769: 1, L0662: 1, L0662: 1, L0776: 1, L0789: 1, L0663: 1, H0658: 1, H0660: 1, L0740: 1, L0758: 1, L0758: 1, L0758: 1, L0758: 1, L0361: 1	L0783: H0662: 3 L0764: 1 L0809: 1	Glu-18 to Asp-35, Leu-52 to Gly-82.					Thr-1 to Ser-8, Pro-22 to Ser-27.
	3424	3425 3426 G	3427	3429 3430	3431	3432	3434 T
·	2 - 289	179 - 541	2 - 454	33 - 425	116-232	3 - 551	1 - 138
	1249	1250	1252 1253	1254 1255	1256	1257 1258	1259
	HNBUY37 HNBUY37R	HNBVL57R HNHBC18R	HNJFE85 HNJFE85R HNKC029 HNKC029R	HNOAA22 HNOAA22R HNOAB27 HNOAB27R	HNOAB28R	HNOAG04 HNOAG04R	HNOAC15 HNOAC15R
	HNBUY37	HNBVL57 HNHBC18	HNJFE85 HNKC029	HNOAA22 HNOAB27	HNOAB28	HNOAE04	HNOAC15

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Thr-5 to Thr-14,	Arg-20 to His-25,	Arg-35 to Gly-40,	Lys-58 to Arg-66,	His-101 to Ser-107,	Arg-111 to Lys-125,	Leu-131 to Glu-140.	Pro-13 to Arg-21,	Ser-42 to Phe-52,	Lys-69 to Val-77,	Ile-82 to Glu-87,	Gly-93 to Gln-99,	Val-118 to Asn-124,	Lys-143 to Leu-152,	Arg-154 to Glu-178.	Thr-S to Thr-14,	Arg-20 to His-25,	Arg-35 to Gly-40,	Lys-58 to Arg-66,	His-101 to Ser-107,	Arg-111 to Lys-125,	Leu-131 to Arg-139,	Arg-145 to Leu-153.	Arg-11 to Pro-17.				Cys-2 to Gly-7.	Trp-6 to Glu-12.
3435					-		3436					-			3437								3438	3439	3440	3441	3442	3443
89 - 577		•					2 - 535								67 - 555				-				2 - 145	42 - 362	28 - 177	2 - 364	3 - 524.	351 - 536
1260				·			1261								1262								1263	1264	1265	1266	1267	1268
HNOAE50 HNOAE50R							HNOAE65R								HNOAF22R								HNOAF36R	HNOAG34R	HNOAG94 HNOAG94R	HNOAH67 HNOAH67R	HNOAJ67R	HNOAJ76R
HNOAE50			·				HNOAE65								HNOAF22								HNOAF36	HNOAG34	HNOAG94	HNOAH67	HNOAJ67	HNOAJ76

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L0748: 7, L0731: 6,	L0591: 6, L0592: 6,	L0770: 5, L0747: 5,	L0749: 5, L0750: 5,	H0136: 5, H0050: 4,	H0625: 4, H0529: 4,	L0769: 4, L0752: 4,	L0758: 4, S0007: 3,	S0045: 3, H0591: 3,	H0561: 3, L0761: 3,	L0645: 3, L0766: 3,	L0655: 3, L0657: 3,	L0666: 3, H0696: 3,	H0436: 3, L0754: 3,	L0756: 3, L0755: 3,	H0543: 3, H0686: 2,	H0125: 2, S0420: 2,	S0376: 2, H0351: 2,	H0581: 2, S0049: 2,	H0052: 2, L0163: 2,	H0428: 2, H0090: 2,	H0040: 2, T0041: 2,	H0494: 2, H0646: 2,	L0763: 2, L0521: 2,	L0776: 2, L0665: 2,	H0144: 2, L0751: 2,	L0745: 2, S0031: 2,	L0581: 2, L0608: 2,	H0170: 1, H0171: 1,	H0394: 1, S0114: 1,
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H0704: 1, S0044: 1, H0187: 1, S0028: 1, L0740: 1, L0779: 1, L0777: 1, L0753: 1, L0759: 1, L0596: 1, L0597: 1, L0595: 1, S0026: 1, H0217: 1 and S0384: 1.					-	,								L0666: 10, H0510: 8,	H0046: 6, H0617: 6,	S0051: 4, H0135: 4,	L0776: 4, L0439: 4,	H0509: 3, L0665: 3,	L0747: 3, H0445: 3,	H0662: 2, S0420: 2,	S0222: 2, H0574: 2,	H0632: 2, H0427: 2,
-				Phe-74 to Pro-81.	Gly-1 to Leu-6,	Pro-47 to Asp-52, Lys-63 to Glu-68	27 5 52 50 50	Arg-9 to Arg-2/,	G19-43 to Lys-46,	Asp-61 to Ser-67,	Ala-124 to Gly-130,	Thr-146 to Lys-151,	Phe-165 to Lys-170.									
·	3444	3445	3446	3447	3448		1	3449						3450				٠				
·	2 - 157	43 - 207	1 - 180	403 - 759	2 - 283			2 - 730						2 - 664								
,	1269	1270	1271	1272	1273			1274						1275								
·	HNOAK11 HNOAK11R	HNOALSIR	HNOAN85R	HNOAO59R	HNOA071R			HNOAPZIK						HNOAQ24 HNOAQ24R								
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H06	H00	H01	H04	106	106	100	H06	L05	106	S02	H02	S03	S60	H02	H05	9 0 0 0	S02)H0	H03	, HO	, HO	, H0	, S60	Ĕ,	H01	H02	, HO	, SO	2
5: 2,	3:2,	5:2,	7:2	0.2,	6:2,	9.2,	1:2,	4.2	9.2,	0:1,	33: 1,	33: 1,	38: 1	11:1	55: 1	31: 1	13: 1	19:1	9:1,	4.	12: 1	24: 1	37:1	56: 1	6: 1	36: 1,	59: 1	47: 1	.0369: 1, L0371
4057	1025	1054	1068	1010	080	.065	590H	L074	1059	3004	H048	H066	H020	H04	H044	H033	H001	H03	S004	H05	H00H	H00.	H01(H02	T000	S003	H00.	H06	8
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	Ser-15 to Glu-23, Val-33 to Gly-40, Arg-47 to Pro-59, Ser-64 to Phe-70, Gly-92 to Lys-97, Lys-155 to Glu-166.	Pro-5 to Gly-11, Ser-20 to Ser-27.
	3451	3452
	1 - 642	122 - 400
·	1276	1277
	HNOAQ47R	HNOAR74R
	HNOAQ47 HNOAQ47	HNOAR74

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H0634: 1, L0637: 1,	L0648: 1, L0649: 1,	L0804: 1, L0774: 1,	L0776: 1, L0655: 1,	L0606: 1, L0809: 1,	H0144: 1, H0698: 1,	L0438: 1, H0690: 1,	H0660: 1, H0651: 1,	L0747: 1, L0749: 1,	L0758: 1, L0601: 1,	S0242: 1, S0196: 1 and	H0543: 1.	L0758: 6, H0031: 4,	H0341: 3, S0022: 3,	S0376: 2, H0586: 2,	H0156: 2, H0575: 2,	L0769: 2, L0803: 2,	L0655: 2, L0809: 2,	H0522: 2, S0027: 2,	H0445: 2, S0412: 2,	H0656: 1, S0358: 1,	S0360: 1, H0574: 1,	H0486: 1, H0013: 1,	H0318: 1, H0327: 1,	L0157: 1, H0015: 1,	H0083: 1, S0003: 1,	S0214: 1, H0039: 1,	H0673: 1, H0591: 1,	T0067: 1, H0412: 1,	H0494: 1, H0529: 1,
												Ser-48 to Leu-56,	Ser-70 to Val-75,	Thr-83 to Ala-92.															-
												3453																_	
												278 - 562))																
												1278																	
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L0372: 1, L0764: 1, L0794: 1, L0649: 1, L0774: 1, L0775: 1, L0651: 1, L0807: 1, L0657: 1, L0789: 1, L0663: 1, H0519: 1, S0126: 1, H0684: 1, H0660: 1, H0666: 1, H0651: 1, H0518: 1, L0777: 1, L0747: 1, S0194: 1, H0543: 1 and S0424: 1.			H0457: 4, H0441: 3, H0393: 2, H0052: 2, H0620: 2, S0144: 2, H0521: 2, H0341: 1, S0358: 1, H0587: 1, H0632: 1, H0581: 1, H0622: 1, H0424: 1, H0213: 1, S0366: 1, H0163: 1, H0132: 1, S0142: 1, H0529: 1,
1.079 1.079 1.065 1.066 8012 8012 8012 1.074 1.074 1.073	Asn-80 to Gly-86, Glu-133 to Arg-138.	Leu-1 to Thr-7, Pro-39 to Ser-48, Trn-62 to Thr-68	* 4. ***********************************
	21 3454	3455	3456
	1279 84 - 521	1280 3 - 254	81 2 - 385
		HNOAS36R 12	HNOAS92R 1281
·	HNOAS07	HNOAS36	HNOAS92

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S3014: 1, L0748: 1,	.0777: 1 and L0599: 1	L0794: 5, H0052: 4	10457: 4, L0438: 4,	H0651: 4, H0581: 3,	J0777: 3, S0116: 2,	S0007: 2, S0222: 2,	S0049: 2, N0006: 2,	J0163: 2, S6028: 2,	J0351: 2, H0130: 2,	J0638: 2, L0803: 2,	J0805: 2, L0439: 2,	.0754: 2, L0750: 2	.0755: 2, H0580:	H0645: 1, H0486: 1,	H0013: 1, S0346: 1	H0009: 1, L0157:		H0050: 1, T0010:	H0606: 1, H0040: 1	H0063: 1, S0210:	L0761: 1, L0667: 1	L0774: 1, L0526: 1	L0532: 1, L0666:	L0665: 1, H0658:	H0521: 1, H0576: 1	L0753: 1, L0758: 1,	L0759: 1, L0593: 1,	L0366: 1, H0543: 1 and	S0462: 1.
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		3457							<u></u>																				
		96 - 317									•••																		
		1282							•																				
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	Lys-28 to Tyr-49,	Arg-75 to Ala-83,	Thr-90 to Ala-95,	Arg-107 to Asp-114,	Arg-137 to Ser-146.		Arg-1 to Gly-6,	Lys-21 to Gly-29,	Gln-51 to Gly-56,	Pro-82 to Asp-90,	Lys-175 to Ser-180.				-													•	
3458	3459					3460	3461																						
2 - 187	2 - 439					98 - 316	2 - 607					•				-							-		-				
1283	1284					1285	1286																_						
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73:3	38: 3	2: 3,	<u>4.</u>	58: 3	10666: 3, S0380: 3,	21:3	H0707: 3, L0589: 3,	0605: 3, L0599: 3,	0593: 3, H0423: 3,	10265: 2, H0661: 2,	S0356: 2, S0358: 2,	S0360: 2, S0408: 2,	S0410; 2, S0046; 2,	H0409: 2, S0280: 2,	10251: 2, L0738: 2,	.0471: 2, H0057: 2,	,0185: 2, H0594: 2,	H0188: 2, S0003: 2,	H0090: 2, H0412: 2,	H0413: 2, S0440: 2,	S0150: 2, S0144: 2,	H0538: 2, L0763: 2,	775:	.0653: 2, H0519: 2,	H0435: 2, H0659: 2,	H0436: 2, H0478: 2,	S0027: 2, L0756: 2,	J0755: 2, L0759: 2,	H0445: 2, L0588: 2,
H03,	H00	2000	7000	90H	S03	H05	165	105	H04	H06	S03;	S04	S 00	. S02	101)H	, но	, SO.	, H0	, SQ	, S01	ë,	5	, HO	2, HO	2, H0	, <u>r</u>	3	2, 1.0
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H0014: 3, H0373: 3	H0553: 3, H0038: 3,	00.	.0761: 3, L0664: 3,	S0126: 3, H0658: 3,	1066	S0152: 3, H0521:	H070	99	1059	H026	S035	S036	S041	H04	H02	1047	<u> </u>	H01	H00	H04	S01.	H05	<u>8</u>	9	HOM	HO HO	80	5	Ž H
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42: 2,	71: 1,	56: 1,	S0342: 1,	35: 1,	, H0671: 1,	49: 1,	25: 1,	34: 1,	15: 1,	70: 1,	22: 1,	40: 1,	35: 1,	56: 1,	10: 1,	10: 1,	46: 1,	93: 1,	10: 1,	67: 1,	52: 1,	39: 1,	81: 1,	68: 1,	40: 1,	67: 1,	64: 1,	25: 1,	41: 1,
L0590: 2, H0542: 2	H0422: 2, H0171:	L0448: 1, H05	H0222: 1, S03	H0650: 1, S00	H0483: 1, H06	•	H0459: 1, H0125:	S0420: 1, S03.	S0444: 1, S00 ²	H0443: 1, H0370: 1	H0392: 1, L06	L0623: 1, T0040: 1	H0013: 1, H0635:	H0427: 1, H0156:	H0590: 1, S0010:	H0052: 1, T0110:	H0597: 1, H0546:	H0009: 1, H0093:		H0510: 1, H0267:	H0687: 1, H0252:	H0428: 1, H0039:	H0622: 1, H0181:	H0032: 1, H0068:		H0087: 1, T0067: 1	-	F0042: 1, H0625:	H0130: 1, H0641:
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																										Pro-14 to Gly-22,	Ile-31 to Ala-36,	Lys-53 to Ile-65.	Val-2 to Leu-20.
									_																	3462			3463
						,			•									•								111 - 506			2 - 394
																										1287			1288
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		H0556: 11, H0265: 9,	H0012: 4, H0616: 4,	H0393: 3, H0620: 3,	L0163: 3, H0135: 3,	S0002: 3, H0651: 3,	L0596: 3, H0542: 3,	H0543: 3, S0418: 2,	S0045: 2, H0266: 2,	S0150: 2, H0521: 2,	L0591: 2, S0040: 1,	H0341: 1, H0663: 1,	S0354: 1, S0358: 1,	S0046: 1, S0222: 1,	H0586: 1, H0486: 1,	H0635: 1, H0618: 1,	٠,	H0263: 1, T0103: 1,		T0003: 1, H0373: 1,	S0051: 1, S0336: 1,	H0290: 1, H0286: 1,	H0328: 1, H0622: 1,	H0553: 1, H0124: 1,	H0090: 1, S0038: 1,	H0100: 1, T0042: 1,	S0015: 1, H0560: 1,		L0654: 1, H0144: 1,
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2 - 193.	1 - 624	1 - 453																										-	
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L0438: 1, H0547: 1, H0519: 1, H0593: 1, H0435: 1, L0602: 1, S0332: 1, H0522: 1, H0134: 1, S3014: 1, L0439: 1, H0445: 1, S0434: 1, L0485: 1, L0604: 1, L0593: 1, L0594: 1, L0601: 1, H0422: 1 and S0398: 1.												S0358: 7, L0750: 7, S0360: 6, H0617: 5, S0126: 4, H0545: 3,
	Pro-1 to Lys-10, Phe-31 to Asn-37, Gln-42 to Leu-50,	Arg-58 to Gln-65,	Leu-86 to Ala-91, Tyr-101 to Ala-111.	Glu-1 to Asp-6.		Ser-47 to Ile-53.	Gly-35 to Glu-41.		Gly-13 to Lys-20, Ala-26 to Lys-32,	Ala-83 to Leu-88, Glv-128 to Val-135,	Phe-142 to Asn-151.	
	3467			3468	3469	3470	3471	3472	3473			3474
	3 - 581			1 - 258	204 - 407	63 - 320	269 - 583	480 - 707	1 - 453	·		2 - 391
	1292	. 		1293	1294	1295	1296	1297	1298			1299
	HNOCR44 HNOCR44R			HNOC1103R	HINODONAL HINODOTAR	HNODG11R	HNODM92 HNODM92R	HNODP28R	HNOJA93R			HNOJB10R
	HNOCR44				HINDCOUS	HNODD//	HNODW92	HNODP28	HNOJA93			HNOJB10

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.0776: 3, S0374: 3	.0748: 3, L.0755: 3,	.0588: 3, H0422: 3	10685: 2, H0549:	0633	942	H0163: 2, H0551: 2,	.0665: 2, H0593: 2,	H0660: 2, S0330: 2,	J0747: 2, L0749: 2,	J0777: 2, L0757: 2,	S0031: 2, H0556:	H0686: 1, H0295: 1	S0430: 1, H0657: 1	H0341: 1, H0662:	S0278: 1, H0282:	H0592: 1, H0586:	H0486: 1, T0039: 1	00	H0581: 1, H0309:	H0596: 1, T0110:	H0231: 1, H0123:	H0620: 1, H0083:	10594: 1, H0687:	.0483: 1, H0644:	40361: 1, H0087:	H0413: 1, H0059:	H0494: 1, S0306: 1	S0440: 1, H0509:	H0529: 1, L0762:
3, S(3, 1,	3, H	2, H	2, H	2, H	2, H	2, H	S. S.	2, Ľ	, , ,	2, H	1, H	1, H	1, H	1, H(1, H	1, T	1, H	1, H	1, T	1, H	1, S(I, H	1,1					
76:	48:	88	385:	50:	.46:	63:	65:	900	47:	77:	31: 3	386:	30:	41:	.8	92:	86:	66	81:	96:	31:	20:	4	83:	61:	.13:	<u>4</u>	6 .	56.
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	Arg-8 to Gln-18, Thr-24 to Gly-35, Lys-53 to Gly-61, Pro-77 to Cys-82, Pro-103 to Gln-117.		Leu-14 to Arg-20, Ser-29 to Ser-38, Pro-43 to Gly-52.	_	Arg-70 to Ser-78.
·	3475	3476	3477	3478	3479
	2 - 355	2 - 439	10 - 165	288 - 572	2 - 283
	1300	1301	1302	1303	1304
	HNOJB57R	HNOJD14R	HNOJF20R	HNO1G05R	HNOJH48R
	HNOJB57	HNOJD14	HNOJF20	TINO 1005	HNOJH48

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L0666: 9, H0547: 8,	L0758: 8, S0358: 7,	S0003: 7, L0754: 7,	L0485: 7, S0280: 6,	H0038: 6, H0521: 6,	S3014: 6, L0777: 6,	L0755: 6, H0068: 5,	L0659: 5, L0665: 5,	L0744: 5, L0740: 5,	L0591: 5, H0657: 4,	H0656: 4, S0116: 4,	H0318: 4, H0266: 4,	H0687: 4, H0031: 4,	H0040: 4, S0002: 4,	S0206: 4, L0752: 4,	H0445: 4, H0542: 4,	H0423: 4, H0497: 3,	H0013: 3, H0427: 3,	H0046: 3, H0014: 3,	H0615: 3, L0142: 3,	S0036: 3, H0090: 3,	H0551: 3, S0440: 3,	L0662: 3, L0767: 3,	H0696: 3, S0027: 3,	S0028: 3, L0731: 3,	L0362: 3, S0196: 3,	H0662: 2, H0638: 2,	H0580: 2, S0045: 2,	S0222: 2, H0587: 2,	H0486: 2, H0706: 2,
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S0010: 2, S0346: 2,	70048: 2, S0474: 2,	10581: 2, H0428: 2,	.0483: 2, H0553: 2,	6: 2, H0634: 2,	H0616: 2, H0412: 2,	H0623: 2, L0763: 2,	i7: 2, L0667: 2,	46: 2, L0768: 2,	74: 2, L0657: 2,	53: 2, H0144: 2,	74: 2, H0520: 2,	H0519: 2, S0126: 2,	28: 2, S0146: 2,	H0555: 2, H0436: 2,	.0747: 2, L0595: 2,	S0026: 2, S0194: 2,	10506: 2, L0600: 2,	70: 1, H0171: 1,	H0556: 1, T0002: 1,	H0686: 1, S0114: 1,	49: 1, S0001: 1,	S0282: 1, H0663: 1,	H0305: 1, S0356: 1,	S0444: 1, H0329: 1,	S0046: 1, L0717: 1,	H1: 1, H0550: 1,	H0441: 1, H0370: 1,	H0600: 1, S0005: 1,	H0333: 1, H0331: 1,
S0010	T004	H058	1.048	8036	H061	H062	L063	1064	1707	99071	8037	H051	8032	H055	L074	8002)50H	HOL	H05	90H	700 <u>T</u>	8028	H03	S04	700S	H04	H04	90H	H03
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H0574: 1. T0082:	H0122-1 H0196-	10122. 1, 110130.	1, 100	0057: 1, H037	010: 1, H051	H0416: 1, S0318: 1,	316: 1, S021	H0074: 1, H0074: 1	H0165: 1, H0168:	H0169: 1, H0674:	S0364: 1, H0708: 1	598: 1, H013	H0163: 1, H0591: 1	H0264: 1, H0272: 1	H0488: 1, H0413:	H0056: 1, S0386:	L0370: 1, T0041: 1	014: 1, H013	H0130: 1, H0647:	S0344: 1, S0422:	.0598: 1, L.0640:	.0770: 1, L0769:	_	J0643: 1, L0771:	.0773: 1, L0766:	•	.0651: 1, L0806: 1		L0655: 1, L0656: 1	.0809: 1, L0790:
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L0791: 1, L0664: 1, H0691: 1, H0689: 1, H0683: 1, H0638: 1, H0660: 1, H0648: 1, H0672: 1, H0539: 1, S0152: 1, S0013: 1, S0174: 1, H0478: 1, L0743: 1, L0750: 1, L0757: 1, S0031: 1, H0707: 1, L0588: 1, L0592: 1, L0599: 1, L0593: 1, H0668: 1, H0667: 1, S0192: 1, S0276: 1, H0422: 1, S0276: 1, H0422: 1,			L0754: 5, L0731: 4, L0748: 3, L0750: 3, H0411: 2, H0012: 2, H0687: 2, H0644: 2, L0803: 2, H0550: 1, L0021: 1, H0599: 1, L0022: 1, H0042: 1,
□ 正正正正ののの日日日日日日 · · · · ·		Lys-36 to Gln-48, Gln-56 to Ser-72, Gly-82 to Val-105, Lys-114 to Lys-120, Ser-122 to Cys-133.	Ala-1 to Ser-6.
	3480	3481	3482
	3 - 200	1 - 489	1 - 552
	1305	1306	1307
	HNOIH52R	HNOJI82R	HNOJI85R
	UNIOIHS		HNOJ185

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H0575: 1, H0150: 1, H0620: 1, T0003: 1, H0673: 1, L0662: 1, L0650: 1, L0774: 1, L0375: 1, L0807: 1, L0515: 1, L0659: 1, L0666: 1, S0374: 1, H0694: 1, L0777: 1 and L0749: 1, L0779: 1 and			L0588: 14, S0046: 13,	S0045: 12, 1'0049: 7, H0599: 7, L0662: 7,	L0599: 7, H0427: 6,	T0048: 6, H0575: 5,	L0747: 5, H0254: 4,	H0618: 4, H0617: 4,	H0124: 4, H0413: 4,	H0052: 3, H0024: 3,	H0553: 3, H0269: 3,	H0412: 3, H0056: 3,	H0623: 3, L0764: 3,	L0659: 3, L0809: 3,	L0666: 3, L0750: 3,	L0758: 3, L0600: 3,	S0001: 2, H0255: 2,	H0661: 2, H0619: 2,
	Lys-19 to Asn-25, Pro-61 to Lys-66.		Ser-27 to Gly-38,	Ser-63 to Trp-72, Trp-79 to Asn-84,	Arg-133 to Trp-139.	•												
	3483	3484	3485							٠.								
	1 - 261	2 - 385	1 - 462								_	٠					_	
	1308	1309	1310															
	HNOJJ20R	HNOJJ26R	HNOJK66R					•							-			
	HNOJJ20	HNOJJ26	HNOJK66															

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S6022: 2, H0587: 2,	S0280: 2, H0253: 2,	H0150: 2, H0050: 2,	H0188: 2, H0644: 2,	S0364: 2, S0036: 2,	.0369: 2, L0768: 2,	2, L0775: 2,	.0375: 2, L0783: 2,	2, L0743: 2,	J0748: 2, L0439: 2,	.0779: 2, L0604: 2,	1, L0785: 1,	1, H0208: 1,	1, H0437: 1,	H0369: 1, H0549: 1,	H0042: 1, H0309: 1,	H0085: 1, H0263: 1,	: 1, S0050: 1,	1, H0266: 1,	H0687: 1, H0039: 1,	: 1, H0030: 1,	H0031: 1, H0163: 1,	H0616: 1, H0063: 1,	: 1, H0268: 1,	.0763: 1, L0536: 1,	: 1, L0638: 1,	0643: 1, L0648: 1,	: 1, L0804: 1,	,0774: 1, L0651: 1,	0378: 1, L0806: 1,
S6022: 2	S0280: 3	H0150:	H0188:	S0364:	10369:	L0803: 2, I	L0375:	L0438: 2, 1	L0748:	L0779:	S6024:	S0360:	L0717:	H0369:	H0042:	H0085:	H0123:	1,0163: 1,	L890H	H0622: 1,	H0031	H0616	H0264: 1,	170763	1,0769: 1,	1.0643	L0794: 1	L0774	1.0378
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HNOJL59	HNOJL59R	1311	204 - 479	3486	Glu-1 to Ser-9, Pro-60 to Gln-66.	50458: 1.		T
HNOJM64	HNOJM64R	1312	3 - 428	3487	Gly-105 to Glu-110, Ala-133 to Tyr-142.			
HNOJO55	HNOJOSSR	1313	426 - 265	3488				Π
HNOJP42	HNOJP42R	1314	12 - 302	3489	Trp-42 to Glu-50.			
HNOJQ22	HNOJQ22R	1315	1 - 510	3490	Phe-164 to Leu-170.		•	
HNOKA20	HNOKA20 HNOKA20R	1316	2 - 268	3491	Gln-30 to Gly-36,			Γ
					Lys-43 to Lys-54,		·	
HNOKD74	HNOKD74 HNOKD74R	1317	2 - 85	3492	TIPLE OF CLASS			T
HNOKG34			1-510	3493	Lys-17 to Leu-27,			T
					Thr-39 to Leu-44,			
					Asp-62 to Gly-70,			
			_		Arg-89 to Asp-94,			
					Arg-102 to Gly-113,			
					Lys-127 to Glu-132,			
					Thr-152 to Arg-160.			
HNOKG69	HNOKG69R	1319	188 - 418	3494				Г
HNOKG71 HNOKG71	HNOKG71R	1320	2 - 331	3495	Gly-8 to Ala-15, Leu-33 to His-50,			<u> </u>
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	L0666: 11, S0358: 8,	L0752: 8, L0749: 5,	L0362: 5, L0771: 4,	L0803: 4, L0528: 4,	L0664: 4, L0740: 4,	L0754: 4, L0599: 4,	L0769: 3, L0662: 3,	L0657: 3, L0659: 3,	L0663: 3, L0748: 3,	L0750: 3, L0757: 3,	L0758: 3, H0171: 2,	S0360: 2, H0574: 2,	H0251: 2, L0483: 2,	H0412: 2, H0509: 2,	L0770: 2, L0774: 2,	L0775: 2, L0665: 2,	H0690: 2, S0330: 2,	L0755: 2, L0731: 2,	L0595: 2, H0624: 1,	H0170: 1, H0685: 1,	H0656: 1, H0663: 1,	H0305: 1, H0589: 1,	S0356: 1, S0354: 1,	H0580: 1, S0132: 1,	S0300: 1, L0717: 1,	H0331: 1, H0013: 1,	H0108: 1, L0105: 1,	H0052: 1, H0546: 1,
Phe-59 to Gly-64,	Gly-70 to Gly-84. Thr-20 to Asp-27,	His-39 to Thr-44.															- '											
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H0123: 1, H0024: 1, H0373: 1, H0266: 1, H0687: 1, L0142: 1, H0628: 1, H0673: 1, H0529: 1, L0763: 1, L0637: 1, L0761: 1, L0646: 1, L0641: 1, L0650: 1, L076: 1, L0650: 1, L0776: 1, H0691: 1, H0519: 1, H0691: 1, H0519: 1, H0648: 1, L0602: 1, S0454: 1, L0439: 1, L0777: 1, L0779: 1, L0779: 1, L0779: 1, L0780: 1, L0753: 1, L0780: 1, L0526: 1, L0759: 1, L0526: 1,	S0194: 1 and H0542: 1. S0410: 13, L0766: 5, H0657: 4, S0422: 4, H0547: 4, S0408: 3, L0758: 3, S0444: 2, H0587: 2, H0486: 2, H0013: 2, H0581: 2, L0471: 2, L0764: 2, L0794: 2, L0666: 2,
	S0194: 1 and H0542 Asp-36 to Glu-46, S0410: 13, L0766: Gln-52 to Ile-61, H0657: 4, S0422: 4, Leu-76 to Lys-87, H0547: 4, S0408: 3, Asn-100 to Val-108, L0758: 3, S0444: 2, Ser-120 to Gln-128. H0587: 2, H0486: 2, H0013: 2, H0581: 2, L0471: 2, L0764: 2, L0794: 2, L0666: 2,
	3497
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L0663: 2, H0520: 2, H0660: 2, H0436: 2,	L0740: 2, L0751: 2,	H0445: 2, S0436: 2,	S0400: 1, H0351: 1,	H0411: 1, H0632: 1,	H0575: 1, S0010: 1,	H0318: 1, H0596: 1,	T0110: 1, H0545: 1,	H0046: 1, H0567: 1,	H0373: 1, S6028: 1,	H0687: 1, H0031: 1,	H0553: 1, H0617: 1,	H0264: 1, S0386: 1,	S0440: 1, H0646: 1,	L0761: 1, L0646: 1,	L0771: 1, L0662: 1,	L0774: 1, L0805: 1,	L0653: 1, L0655: 1,	L0664: 1, H0519: 1,	H0521: 1, H0478: 1,	L0756: 1, L0752: 1,	L0759: 1, S0434: 1,	L0362: 1, S0192: 1 and	H0543: 1.			L0751: 9, H0556: 8,	H0617: 8, L0777: 6,
					-																			Glu-1 to Pro-7,	Asp-25 to Leu-31, Arg-54 to Lvs-73.	Pro-7 to Gly-20.	
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.0809: 5, L0741: 5,	.0754: 5, L0779: 5,	S0222: 4, L0769: 4,	.0659: 4, L0438: 4	.0748: 4, H0208: 3,	H0052: 3, H0009: 3,	30051: 3, T0010: 3,	H0521: 3, L0747: 3,	JO731: 3, LO758: 3,	S0408: 2, S0476: 2,	H0575: 2, H0253: 2,	H0545: 2, H0687: 2,	H0615: 2, H0181: 2,	H0135: 2, H0494: 2,	.5566: 2, L0666: 2,	.0664: 2, L0565: 2	H0445: 2, S6024: 1,	S0134: 1, S0001: 1,	S0282: 1, S0400: 1	0354:	S0360: 1, H0580:	S0007: 1, H0639: 1	H0441:	H0370: 1, H0438: 1,	30280: 1, H0036:	H0618: 1, S0049:	H0196: 1, H0235: 1	H0597: 1, H0231: 1	H0327: 1, H0566: 1,	.0163:
9: 5, L	4: 5, L	2: 4, L	9: 4, L	8: 4, H	52: 3, E	1:3, T	11: 3, L	1: 3, L	8: 2, S	75: 2, E	15: 2, F	15: 2, F	5: 2, F	6: 2, L	4: 2, L	15: 2, S	4: 1, S	2: 1, S	39: 1, S	0: 1, H	7: 1, H	S0278: 1, H	70: 1, E	0: 1, H	8: 1, S	6: 1, E	77: 1, E	7: 1, E	H0024: 1, L0163:
1080	1.075	S022	1065	L074	H005	S005	H052	L073	S040	H057	H054	H061	H013	L556	1066	H04	S013	S028	H045	S036	<u>2000</u>	S027	H037	S028	H061	H019	H059	H032	H002
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.0756: 5, L0731: 5,	0766: 4, L0748: 4,	S0212: 3, L0598: 3,	0637: 3, L0761: 3,	0664: 3, H0658: 3,	1,104	, L07,	0755: 3, L0758: 3,	.0604: 3, L0595: 3,	H0686: 2, L0002: 2,	S0376: 2, S0360: 2,	S6026: 2, H0351: 2,	S0222: 2, H0455: 2,	.0471: 2, H0014: 2,	10006: 2, S0440: 2,	S0422: 2, L0667: 2,	.0641: 2, L0764: 2,	, L07,	, LO52	, 1.078	S0374: 2, H0539:	, L055				H0170: 1, S0218: 1	H0657: 1, H0656: 1	L0808: 1, S0400: 1	H0402: 1, S0442: 1,	S0444: 1, S0408: 1
756: 5	766: 4	212: 3	637: 3	664: 3	0742: 3,	0747: 3,	755: 3	604:3	686: 2	376: 2	256: 2	222: 2	471:2	306: 2	122: 2	541:2	.0768: 2, 1	.0776: 2, 1	.0636: 2, 1	374: 2,	.0749: 2,]	.0589: 2, 1	.0592: 2,]	34: 2	170:1	557: 1	308: 1,	402: 1	144: 1,
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H0441: 1	H0415: 1, H0587: 1	H0486	r0039: 1, H0156: 1,	H0318: 1, H0173:	H059	L0041: 1, H0011: 1,	H037	S005	H039	H0061: 1, H0687: 1	L005	H055	T004	SK	1076	T076	, L0521: 1,	7,07	1065	L0653:]	10907	L0512:	2,2	L0517: 1, L0526: 1,	7,	.2269: 1, L0438:	HO519: 1, H0659:	H0660: 1, H0651: 1	S0328: 1, S0380: 1,
<u>o</u>	; -	2.	9: 1,	18: 1,	96: 1,	11: 1,	57: 1,	51: 1,	10: 1,	61: 1,	28: 1,	73: 1,	13: 1,	49: 1,	.0762: 1,	_	_	0364: 1,		.0378: 1,	,0606: 1,	0661: 1,	56: 1	17:1	309: 1	269: 1	519: 1	660: 1	328: 1
H0549-1	104	1.062	1003	H03	HOI	<u>Š</u>	<u>8</u>	0H	100	HOO	H03	90H	H04	90H	107	2	<u>17</u>	<u>1</u>	<u>~</u>	<u>8</u>	<u> </u>	<u> </u>	<u> </u>	Š	<u> </u>	17	Ë	윤	SO
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H0710: 1, H0579: 1, S0404: 1, S0406: 1, L0740: 1, L0754: 1, L0777: 1, L0759: 1, S0308: 1, L0605: 1, L0581: 1, L0608: 1, L0601: 1, H0543: 1, S0412: 1, S0424: 1 and H0506: 1.				H0624: 10, S0192: 8, S0022: 7, S0212: 5, H0031: 4, H0412: 4, S0028: 4, L0747: 4, S0194: 4, H0717: 3, H0427: 3, L0471: 3, S0250: 3, H0039: 3,
	Lys-1 to Glu-6, Pro-8 to Gln-21, Ser-43 to Glu-51, Val-61 to Gly-68, Arg-75 to Pro-87, Ser-92 to Phe-98.	Lys-13 to Ser-41, Lys-57 to Lys-66, Lys-89 to Lys-97, Lys-113 to Ala-118, Pro-131 to Ala-147, Ala-159 to Ala-169.	Lys-24 to Lys-31, Gly-132 to Glu-137.	
	3501	3502	3503	3504
,	98 - 457	3 - 509	3 - 488	216 - 401
	1326	1327	1328	1329
·	HŅOKT24R	HNOKUS2 HNOKUS2R	HNOKV82R	HNOKZ03R
,	HNOKT24	HNOKU52	HNOKV82	HNOKZ03 HNOKZ03

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H0644: 3, H0040: 3,	H0059: 3, H0144: 3,	H0547: 3, S0126: 3,	80037: 3, 80027: 3,	\$0206: 3, L0439: 3,	H0713: 2, H0661: 2,	S0360: 2, H0208: 2,	S0045: 2, S0046: 2,	H0587: 2, H0486: 2,	H0575: 2, H0024: 2,	H0266: 2, H0622: 2,	H0551: 2, S0210: 2,	H0520: 2, L0740: 2,	.0731: 2, L0605: 2,	H0170: 1, H0171: 1,	S0040: 1, H0716: 1,	H0294: 1, S0116: 1,	S0420: 1, S0442: 1,	0619: 1, S0300: 1,	0592: 1, H0586: 1,	H0485: 1, H0156: 1,	H0590: 1, S0474: 1,	S0049: 1, H0596: 1,	0544: 1, H0045: 1,	H0012: 1, H0620: 1,	H0014: 1, T0010: 1,	H0687: 1, S0003: 1,	H0615: 1, H0030: 1,	S0364: 1, S0366: 1,	H0090: 1, H0038: 1,
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H0272: 1, H0334: 1, S0150: 1, L0598: 1, L0767: 1, L0766: 1, L0791: 1, L0792: 1, L0665: 1, T0068: 1, L0438: 1, H0660: 1, S3014: 1, L0744: 1, L0754: 1, L0749: 1, L0757: 1, L0759: 1, L0588: 1, L0589: 1, L0591: 1, L0592: 1, S0011: 1, H0665: 1,	·		S0412: 22, L0662: 19, L0777: 14, S0010: 11, S0222: 9, S6028: 9, L0750: 9, T0010: 8, L0659: 8, L0747: 8, L0756: 8, L0663: 7, L0439: 7, H0051: 6, L0518: 6, L0754: 6, L0752: 6, S0280: 5,
	Leu-24 to Thr-39, Gln-70 to Glu-81, Arg-86 to Leu-98, Glu-104 to Pro-110, Pro-134 to Arg-157, Val-159 to Asn-166.	Gly-1 to Ser-22, Pro-29 to Lys-70.	
·	3505	3506	3507
	2 - 595	1 - 210	2 - 499
	1330	1331	1332
	HNORAS6 HNORAS6R	HNORB68 HNORB68R	HNORC14R
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358: 4,	753: 4,	3, 3,	427: 3,	373: 3,	L0637: 3,	L0774: 3,	666: 3,	672: 3,	731:3,	0713: 2,	411:2,)574: 2,	486: 2,	049: 2,	327: 2,	0428: 2,	3644: 2,)520: 2,	770: 2,	517: 2,	L0442: 1,)443: 1,	282: 1,	661: 1,	356: 1,	329: 1,	619: 1,	300: 1,	022: 1,
H0575: 5, S0358: 4	S0346: 4, L0753: 4,	H0170: 3, H0662: 3,	S0360: 3, H0427: 3,	.0471: 3, H0373: 3,	.0638: 3, LO	.0764: 3, LO	J0809: 3, L0666: 3,	S0310: 3, H0672: 3,	H0696: 3, L0731: 3,	HO506: 3, H0713: 2,	S0408: 2, H0411: 2,	H0455: 2, H0574: 2,	S0414: 2, H0486: 2,	H0036: 2, S0049: 2,	H0052: 2, H0327: 2,	H0687: 2, H0428: 2,	H0553: 2, H0644: 2,	H0038: 2, L0520: 2,	.0762: 2, L0770: 2,	J0653: 2, L0517: 2	30044: 2, LO	H0583: 1, L0443: 1	S0110: 1, S0282:	S0400: 1, H0661:	S0348: 1, S0356:	S0444: 1, H0329: 1	S0007: 1, H0619: 1,	S6026: 1, S0300: 1,	.0717: 1, S6022:
H05	S034	H01,	803	107	1063	107	1080	S031	90H	H05(S040	H04	S041	H00.	H00	90H	HOS	HOOH	107	106	800	HOS	S011	S04(S03	S04	2000	S602	L07
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, Н0592:	H0587: 1, H0599:	H0042: 1, H0590:	1, H0309:	H0545: 1, H0563:	H0564: 1, H0123:	H0019: 1, S0050:	, S0388:	H0356: 1, H0328:	10688: 1, H0039:	C0023: 1, H0031:	H0111: 1, H0169:	S0364: 1, L0455:	H0135: 1, H0163:	F0067: 1, T0069:	H0100:	, L0370:	T0769:	L0800: 1	L0363: 1	L0649:	L0775:	, L0651:	1, L0523:	L0776:	.0527: 1, L0657:	L0783: 1	053	H0691:]	F0068: 1, S0148: 1
1,1	1, F	1, F	1, F	1, E	1, F	1, S	ı, S	1,1	1, E	Ξ̈́	1,		H. H.	. T	Œ,			^			- :		Ľ,	•	1	ij		耳	š
ő	87:	#	18:	45:	4.	61		56:	.: 88:	23	11:	4.	35:		4	2	 	ö	<u></u>	 	33:]	5: 1	4.	5:]			9	4.	
H0550: 1	105	<u>8</u>	10318:	105	1050	8	.0163: 1	103	90	90	101	36	101	ğ	T0004: 1	S0112:	10598: 1	.0630: 1	.0648:]	.0768: 1	0803: 1,	.0375: 1	.0784: 1	0805: 1	92	.0635: 1,	.0789: 1,	L0664: 1	8
-	يلز	<u> </u>	<u> 11</u>	<u>;11;</u>	<u>ili</u>	<u>;11;</u>		<u>11</u>	<u>. 11 i</u>	Н	يلز	S	نكز	=	H	<u> </u>	<u></u>	브	<u> </u>	ᆜ	<u>, , , , , , , , , , , , , , , , , , , </u>			ㅂ			그	<u>,,</u>	_
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H0693: 1, H0520: 1, H0593: 1, H0689: 1, H0684: 1, S0330: 1, S0380: 1, S0174: 1, H0555: 1, L0612: 1, L0743: 1, L0748: 1, L0751: 1, L0749: 1, L0779: 1, L0759: 1, L0689: 1, S0434: 1, L0604: 1, L0366: 1, S0106: 1 and S0021: 1.		H0585: 20, H0641: 14, H0521: 9, H0141: 5, H0031: 5, H0422: 5, H0644: 4, L0755: 4, H0295: 3, H0638: 3, S0340: 3, H0689: 3, H0689: 3, H0689: 3, L0757: 2, S0360: 2, H0657: 2, S0360: 2, H0662: 2, H0691: 2, H0522: 2, L0740: 2, L0777: 2, H0423: 2, S0298: 1, H0306: 1, S0228: 1, H0549: 1, S0222: 1, H0549: 1, S0222: 1,
		Ala-24 to Lys-29, Lys-51 to Gly-62.
	3508	3509
	155 - 430	2 - 436
	1333	
	HNORD11R	HNOREGSR
·	HNORD11	HNORE65

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H0587: 1, H0333: 1, H0575: 1, H0706: 1, H0596: 1, H0546: 1, H0596: 1, H0546: 1, H0545: 1, H0284: 1, T0023: 1, L0143: 1, H0551: 1, H0494: 1, S0438: 1, S0002: 1, L0371: 1, L0770: 1, L0372: 1, L0646: 1, L0372: 1, L0646: 1, L0774: 1, L0382: 1, H0682: 1, S0328: 1, S0380: 1, H0593: 1, H0682: 1, S0328: 1, S0380: 1, H0555: 1, L0747: 1, L0779: 1, L0747: 1, L0779: 1, L0747: 1, L0779: 1,	L0439: 6, L0794: 2, L0665: 2, H0691: 2, L0717: 1, H0068: 1, L0761: 1, L0646: 1, L0773: 1, L0662: 1, L0790: 1, L0663: 1, L0438: 1, L0748: 1, L0758: 1, L0592: 1 and
	Pro-19 to His-36, Glu-68 to Ala-86.
	3510
	37 - 387
	1335
	HNORF03R
	HNORF03

L0608: 1.				S0358: 7, L0750: 7,	S0126: 4, H0545: 3,	L0776: 3, S0374: 3,	L0748: 3, L0755: 3,	L0588: 3, H0422: 3,	H0685: 2, H0549: 2,	H0550: 2, H0632: 2,	H0546: 2, H0424: 2,	H0163: 2, H0551: 2,	L0665: 2, H0593: 2,	H0660: 2, S0330: 2,	L0747: 2, L0749: 2,	L0777: 2, L0757: 2,	S0031: 2, H0556: 1,	H0686: 1, H0295: 1,	S0430: 1, H0657: 1,	H0341: 1, H0662: 1,	S0278: 1, H0282: 1,	H0592: 1, H0586: 1,	H0486: 1, T0039: 1,	T0109: 1, H0004: 1,	H0581: 1, H0309: 1,	H0596: 1, T0110: 1,
		Thr-2 to Phe-7, Gly-18 to Arg-23, A sp-27 to Glu-35	Gly-1 to Thr-10.								-															
	3511	3512	3513	3514					-																	
	1 - 168	2 - 334	1-636	3 - 680				-					·												•	
	1336	1337	1338	1339																						
	HNORF04R	HNORF05R	HNORF13R	HNORH03R																						
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H0231: 1, H0123: 1, H0520: 1, H0083: 1, H0594: 1, H0687: 1, L0483: 1, H0687: 1, H0361: 1, H0087: 1, H0494: 1, S0306: 1, S0440: 1, H0509: 1, L0770: 1, L0762: 1, L0770: 1, L0769: 1, L0771: 1, L0773: 1, L0771: 1, L0773: 1, L0775: 1, L0375: 1, L0807: 1, L0656: 1, L0659: 1, L0656: 1, H0691: 1, H0144: 1, S0216: 1, H0144: 1, S0216: 1, H0144: 1, H0691: 1, H0678: 1, H0691: 1, H0678: 1, H0684: 1, H0678: 1, S0328: 1, S0378: 1, S0328: 1, S0378: 1, S0328: 1, L0741: 1, H0445: 1, L0596: 1,	S0460: 1 and H0352: 1.	
	Arg-13 to Arg-21,	Asp-26 to Pro-35,
	3515	
	1 - 498	
	1340	
	HNORH05 HNORH05R	
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												L0742: 10, H0144: 6,	H0265: 5, S0116: 4	S0046: 4, S0222: 4	H0556: 3, H0250: 3,	H0052: 3, S0027: 3	L0595: 3, S0212: 2,	H0306: 2, S0358: 2	H0013: 2, H0618:	H0124: 2, H0100: 2,	S0053: 2, H0660: 2,	S0330: 2, S3012: 7	L0748: 2, L0439:	L0745: 2, L0757: 2,	H0445: 2. S0342:	H0484: 1, H0638: 1	S0007: 1, H0393:	H0437: 1, H0549: 1,	
Glu-58 to Gly-65,	Ser-74 to Pro-82,	Gln-107 to Asp-115,	Asp-128 to Asn-	137,	Gln-145 to Arg-154,	Gln-161 to Ser-166.	Ala-19 to Ser-27,	Ser-41 to Val-53.	Asn-95 to Leu-100.	Pro-3 to Ile-12,	Asn-20 to Glu-25.	He-121 to Val-129,	Ala-136 to Glu-142. H0265: 5, S0116: 4,												·				
							3516		3517	Γ		3519																	
							20 - 244	•	3 - 377	144 - 377	· ·	3.530	5																
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ł	H0253: 1, H0390:	÷.	_	N0007: 1, T0010:	÷,	S0003: 1, H0030:	S0364: 1, L0455:	S0366: 1, H0090:	H0038: 1, H0551:	H0412: 1, H0059:	S0038: 1, H0130:	H0641: 1, S0142:	S0002: 1, H0517:	<u>-</u>	L0794: 1, L0766:	L0775: 1, L0654:	L0655: 1, L0352:	S0008: 1, H0435:	H0659: 1, H0658:	H0672: 1, H0521:	H0522: 1, S0044:	H0576: 1, L0779:	L0605: 1, L0593:	S0026: 1, H0667:	S0276: 1, S0196: 1	H0542: 1, H0423: 1 and	H0506: 1.	L0439: 17, L0438: 6,	H0424: 5, S0007: 4, I 0769: 4 H0213: 3	
								•							•	3												Asn-39 to Gln-44,	Leu-47 to Asp-55.	
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L0776: 3, H0660: 3, H0309: 2, S0051: 2, L0789: 2, L0663: 2, L0759: 2, L0663: 2, L0759: 2, L0757: 2, L0759: 2, S0196: 2, S0040: 1, S001: 1, H0662: 1, H0402: 1, H0052: 1, H0596: 1, H0009: 1, T0010: 1, L0483: 1, L006: 1, H0538: 1, L0764: 1, L0643: 1, L0764: 1, L0663: 1, L0764: 1, L0791: 1, L0375: 1, L0791: 1, L0372: 1, L0791: 1, L0372: 1, L0752: 1, L0777: 1, L0752: 1, L0772: 1, L0758: 1, S0394: 1, L0758: 1, S0394: 1,		·
		Lys-18 to Tyr-31, Ile-45 to Asp-50, Trp-78 to Ser-88,
	3521	3522
	241 - 456	1 - 426
	1346	1347
	HOCMB65 HOCMB65R	HOCMC08 HOCMC08R
	HOCMB65	HOCMC08

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			<u> </u>	L0803: 7, H0169: 6,	H0144: 6, L0731: 4,	H0574: 3, H0521: 3,	H0370: 2, H0615: 2,	H0032: 2, T0067: 2,	H0509: 2, S0210: 2,	L0776: 2, L0749: 2,	L0750: 2, L0595: 2,	H0170: 1, H0656: 1,	H0341: 1, H0151: 1,	S0046: 1, S0132: 1,	H0619: 1, H0351: 1,	S0222: 1, H0497: 1,	H0632: 1, H0486: 1,	H0156: 1, H0575: 1,	H0052: 1, H0596: 1,	H0057: 1, H0014: 1,	H0355: 1, H0286: 1,	H0622: 1, H0165: 1,	H0674: 1, H0212: 1,	H0038: 1, H0379: 1,	H0059: 1, T0041: 1,	S0448: 1, H0132: 1,	S0150: 1, L0646: 1,	L0774: 1, L0527: 1,	L0657: 1, L0809: 1,
Pro-102 to Pro-109.		Glu-1 to Ile-6,	Asp-125 to Pro-131.			-																				•			
	3523	3524		:																									
	382 - 588	3 - 395											•						,										
	1348	1349																											
	HOCMF06 HOCMF06R	HOCMF27 HOCMF27R																											
	HOCMF06	HOCMF27				,																	7						

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L0519: 1, L0788: 1, L0789: 1, L0666: 1, H0520: 1, H0519: 1, H0658: 1, H0660: 1, S0152: 1, S0013: 1, H0696: 1, S0044: 1, S0174: 1, S0146: 1, H0479: 1, L0744: 1, L0777: 1, L0758: 1, L0581: 1, L0608: 1,	H0687: 1, L0663: 1 and H0660: 1.	L0794: 7, L0666: 4, L0750: 3, L0664: 2, L0665: 2, H0670: 2, H0660: 2, L0439: 2, L0751: 2, L0756: 2, H0650: 1, H0150: 1, L0769: 1, L0803: 1, L0378: 1, L0657: 1, H0684: 1 and H0555: 1,	H0657: 13, S0360: 12, S0026: 7, H0547: 6, H0171: 4, S0356: 4, H0551: 4, H0560: 4, H0599: 4, H0599: 4, H0543: 4, S0358: 3,
	Gly-2 to Lys-9, Arg-21 to Glu-32, Pro-58 to Lys-64, Thr-73 to Glu-83.	Lys-1 to Thr-6, Ala-26 to Gly-36, Ser-47 to Gln-55, Leu-72 to Cys-83.	Gly-1 to Asn-9, Glu-31 to Val-39.
	3525	3526	3527
	49 - 354	37 - 285	1 - 456
	1350	1351	1352
	HOCMF50 HOCMF50RB	HOCMF54R	HOCMG37R
	HOCMF50	HOCMF54	HOCMG37

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H0318: 3, S0440: 3, H0529: 3, H0435: 3,	H0665: 3, H0170: 2, H0265: 2, H0556: 2	10222: 2, 50376: 2,	.0717: 2, H0642: 2,	H0574: 2, H0575: 2,	H0596: 2, S0003: 2,	H0615: 2, H0428: 2,	H0617: 2, H0413: 2,	10494: 2, H0646: 2,	.0369: 2, H0144: 2,	, H0519: 2,	10690: 2, H0658: 2,	10660: 2, H0672: 2,	,0362: 2, H0542: 2,	10423: 2, H0686: 1,	H0650: 1, H0656: 1,	S0212: 1, S0029: 1,	H0483: 1, H0663: 1,	H0664: 1, H0638: 1,	S0420: 1, H0580: 1,	S0278: 1, H0261: 1,	, H0586: 1,	roo60: 1, H0069: 1,	S0010: 1, H0263: 1,	H0327: 1, H0546: 1,	H0014: 1, H0083: 1,	H0510: 1, S6028: 1,	H0266: 1, S0214: 1,
H0318: 3 H0529: 3	H0665: 3	H0222: 2	L0717: 2	H0574: 2	H0596: 2	H0615: 2	H0617: 2	H0494: 2	L0369: 2	H0520: 2	H0690: 2	H0660: 2	1.0362: 2	H0423: 2	H0650: 1	S0212: 1	H0483: 1	H0664: 1	S0420: 1	S0278: 1	H0438: 1	T0060: 1	S0010: 1,	H0327: 1	H0014: 1	H0510: 1	H0266: 1
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H0553: 1, H0068: 1, H0090: 1, H0040: 1, H0634: 1, T0067: 1, H0079: 1, H0561: 1, L0065: 1, H0130: 1, L0772: 1, L0648: 1, L0659: 1, L0768: 1, L0659: 1, L0518: 1, H0670: 1, H0648: 1, S0310: 1, L0438: 1, H0670: 1, H0648: 1, L0749: 1, L0738: 1, L0779: 1, L0731: 1, H0343: 1, L0596: 1, L0591: 1, L0603: 1, H0653: 1, S0424: 1, H0653: 1, S0424: 1, S0384: 1 and H0506: 1.					L0748: 2, S6028: 1, H0682: 1, H0660: 1,
		Lys-23 to Ser-29, Tyr-55 to His-62, Tyr-65 to Leu-71.	Glu-32 to Ser-38.	Gly-1 to Arg-9.	
	3528	3529	3530	3531	3532
	49 - 348	1 - 510	14 - 400	3 - 707	2-289
	1353	1354	1355	1356	1357
			HOCMK20R	HOCMI OFR	HOCML66R
	HOCMI62	HOCMJ47	HOCMMESO	HOCKIT OF	HOCML66 HOCML66R

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S0378: 1 and H0506: 1.			H0295: 8, H0617: 7,	L0665: 7, H0435: 7,	H0592: 6, H0494: 6,	L0439: 6, L0751: 6,	S0360: 5, S0046: 4,	H0587: 4, H0012: 4,	H0264: 4, L0662: 4,	S0406: 4, H0586: 3,	H0620: 3, H0031: 3,	H0551: 3, S0440: 3,	S0002: 3, L0763: 3,	L0648: 3, L0649: 3,	L0774: 3, L0659: 3,	H0520: 3, H0547: 3,	H0689: 3, H0660: 3,	S0328: 3, S0330: 3,	L0748: 3, L0754: 3,	L0750: 3, H0294: 2,	H0650: 2, H0341: 2,	80356: 2, 80132: 2,	H0370: 2, H0333: 2,	H0643: 2, L0623: 2,	H0706: 2, H0081: 2,	H0039: 2, H0553: 2,	H0181: 2, H0068: 2,	H0135: 2, H0040: 2,
	Thr-19 to Cys-25, Lys-133 to Ala-138.																					<u> </u>					H-4.	
	3533	3534	3535																									-
	130 - 594	3 - 425	3 - 500																				-				,	
	1358	1359	1360															-										
	HOCMM52 HOCMM52R	HOCMO50 HOCMO50R						•					•	-		_							-			_		
	HOCMM52	HOCMO50	HOCMS75								-										-		-					

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H0646: 2, L0640: 2,	.0637: 2, L0641: 2,	.0764: 2, L0804: 2,	.0806: 2, L0776: 2,	20657: 2, L0809: 2,	.0664: 2, H0519: 2,	H0682: 2, H0670: 2,	H0672: 2, L0744: 2,	.0740: 2, S0434: 2,	H0677: 2, H0483: 1,	H0305: 1, H0638: 1,	S0442: 1, S0358: 1,	S0376: 1, S0410: 1,	H0411: 1, H0550: 1,	H0455: 1, H0600: 1,	H0101: 1, H0575: 1,	H0590: 1, H0618: 1,	H0052: 1, H0234: 1,	H0597: 1, H0121: 1,	L0041: 1, H0123: 1,	H0246: 1, H0290: 1,	H0292: 1, H0428: 1,	H0622: 1, T0023: 1,	H0030: 1, H0316: 1,	H0059: 1, H0100: 1,	S0306: 1, T0090: 1,	H0649: 1, S0210: 1,	S0426: 1, L0646: 1,	L0765: 1, L0771: 1,	L0626: 1, L0768: 1,
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L0364: 1, L0766: 1, L0375: 1, L0651: 1, L0378: 1, L0653: 1, L0655: 1, L0558: 1, L0635: 1, L0578: 1, L0383: 1, L0792: 1, L0663: 1, S0006: 1, H0696: 1, S0044: 1, S3014: 1, L0743: 1, L0752: 1, L0755: 1, L0751: 1, L0759: 1, H0216: 1, H0543: 1,	L0743: 9, H0551: 5, H0645: 4, H0050: 4, S0250: 4, S0192: 4, H0624: 3, T0039: 3, H0087: 3, H0294: 2, H0620: 2, H0024: 2, H0622: 2, H0424: 2, H0622: 2, H0424: 2, H0622: 2, L0776: 2, S0028: 2, L0776: 2, S0242: 2, S0116: 1, H0484: 1, H0483: 1, S0360: 1, H0619: 1, H0550: 1, H0619: 1,
	Pro-84 to Pro-91.
	3536
	145 - 636
	1361
	HOCMS87R
	HOCMS87

H0599: 1, H0575: 1, H0122: 1, H0706: 1, H0421: 1, S0049: 1, H0086: 1, H0123: 1, H0081: 1, H0012: 1, H0687: 1, H0028: 1, H0604: 1, S0386: 1, L0772: 1, L0748: 1, L0375: 1, L0774: 1, L0375: 1, L0778: 1, H0684: 1, H0660: 1, S0152: 1, S0027: 1, L0779: 1, L0749: 1, L0779: 1, L0777: 1,							•	L0777: 14, L0731: 10, L0740: 8, L0439: 7, L0754: 6, S0222: 5, H0144: 5, L0745: 5,
		Glu-1 to Gly-6, Leu-27 to Val-40,	Glu-60 to Asn-67.		Gly-17 to Arg-42, Phe-56 to Ser-66.		Lys-1 to His-6.	Lys-7 to Glu-13, Gly-39 to Thr-44, Gly-82 to Ile-89, Glu-98 to Asn-105.
	3537	3538		3539	3540	3541	3542	3543
	86 - 235	76 - 354		2 - 466	314-511	394-2	520 - 789	3 - 488
	1362	1363		1364	1365	1366	1367	1368
	HOCMX29 HOCMX29R	HOCMY53 HOCMY53R		HOCMY61R	HOCMY79 HOCMY79R	HOCOC38R	HOCOC41R	HOCOC94R
· · · · · · · · · · · · · · · · · · ·	HOCMX29	HOCMY53		HOCMY61	HOCMY79	HOCOC38	HOCOC41	HOCOC94

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H0013: 4, L0770: 4,	S0380: 4, L0748: 4,	L0758: 4, H0171: 3,	H0599: 3, S0010: 3,	H0052: 3, S0003: 3,	H0615: 3, L0598: 3,	L0666: 3, L0665: 3,	H0519: 3, H0658: 3,	L0749: 3, H0255: 2,	S0376: 2, S0444: 2,	S0360: 2, H0632: 2,	H0327: 2, S0214: 2,	H0038: 2, H0040: 2,	L0769: 2, L0638: 2,	L0662: 2, L0794: 2,	L0775: 2, L0776: 2,	H0660: 2, S0330: 2,	L0744: 2, L0750: 2,	L0756: 2, L0757: 2,	L0362: 2, L0361: 2,	H0170: 1, H0685: 1,	S0040: 1, S0342: 1,	H0716: 1, S0001: 1,	S0282: 1, H0662: 1,	S0356: 1, S0358: 1,	H0675: 1, S0408: 1,	H0580: 1, S0300: 1,	H0406: 1, S0278: 1,	H0550: 1, H0441: 1,	H0431: 1, H0600: 1,
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H0331: 1, H0574: 1.	r0039: 1, H0156: 1,	S0665: 1, S0346: 1,	H0263: 1, H0596: 1,	10546: 1, H0150: 1,	I0009: 1, H0569: 1,	L0471: 1, H0012: 1,	H0051: 1, S0051: 1,	10010: 1, H0266: 1,	30250: 1, S0022: 1,	10428: 1, H0119: 1,	10553: 1, H0628: 1,	H0617: 1, H0316: 1,	10634: 1, H0412: 1,	H0413: 1, H0623: 1,	H0494: 1, S0450: 1,	H0130: 1, S0142: 1,	UNKWN: 1, L0771: 1,	L0766: 1, L0803: 1,	L0804: 1, L0774: 1,	L0784: 1, L0805: 1,	L0652: 1, L0653: 1,	L0655: 1, L0606: 1,	L0527: 1, L0783: 1,	L0663: 1, H0547: 1,	H0689: 1, H0670: 1,	H0648: 1, H0539: 1,	H0521: 1, H0696: 1,	H0478: 1, S0432: 1,	H0540: 1, S0028: 1,
<u> </u>	Ē	<u> </u>	正	<u>=</u>	正	ı	<u>:</u>	<u>.</u>	· ·	<u>;1;</u>	<u>. i.l.</u> i	14		<u> </u>				- - 											
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S0206: 1, L0746: 1, L0747: 1, S0031: 1, L0689: 1, S0436: 1, S0194: 1, S0276: 1, S0412: 1, H0506: 1 and H0352: 1.					H0521: 5, S0040: 3, H0039: 3, H0657: 2, H0341: 2, H0553: 2, S0210: 2, H0593: 2, H0445: 2, L0596: 2, S0212: 1, H0661: 1, S0418: 1, S0354: 1, S0358: 1, H0592: 1,
· .	Pro-6 to Arg-12, Gln-17 to Leu-22, His-25 to Asp-32, Gly-62 to Lys-69.	Thr-16 to Phe-28, Cys-58 to Ser-69, Glu-91 to Glu-99, Lys-113 to Gln-119.	Gin-37 to Trp-44, Arg-47 to Phe-57, Phe-67 to Ala-83, Pro-94 to Gly-99, Lys-109 to Cys-116, Arg-130 to Ser-137.		Glu-1 to Val-9.
	3544	3545	3546	3547	3548
	221 - 502	2 - 409	9 - 539	2 - 382	46 - 492
	1369	1370	1371	1372	1373
	HOCOF35R	HOCOF50R	носоо62к	HOCOP52R	носоотзя
	HOCOF35	HOCOFSO	НОСОО62	HOCOP52	носоо13

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H0427: 1, H0544: 1, H0546: 1, H0545: 1, H0046: 1, L0471: 1, H0083: 1, S6028: 1, S0338: 1, S6028: 1, H0068: 1, H0038: 1, H0068: 1, H0551: 1, H0494: 1, H0551: 1, H0649: 1, L0654: 1, L0666: 1, H0660: 1, H0518: 1, S0152: 1, S0146: 1, H0555: 1, S0146: 1, H0555: 1, L0605: 1, L0599: 1, L0605: 1, H0543: 1, L0605: 1, H0543: 1, R0542: 1, H0543: 1,					H0046: 5, H0144: 5, .0665: 4, L0662: 3,	,0766: 3, L0750: 3,	0779: 3, L0759: 3,	H0170: 2, S0376: 2,	H0329: 2, H0427: 2,
<u> </u>			Lys-9 to Asp-16, Thr-27 to Trp-36.		3553 Pro-38 to Gln-43.	<u> </u>	<u>-</u>	<u></u>	
·	3549	3550	3551	3552	3553			-	
	448 - 639	2-412	107 - 268	1 - 573	1 - 183				
	1374	1375	1376	1377	1378				
	HOCOQ93R	HOCOT70R	HOCOV29R	HOCPF69R	HOCPG38R	-			
	HOC0093	HOCOT70	HOCOV29	HOCPF69	HOCPG38				

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.0770: 2, L0529: 2,	0439: 2, L0740: 2,	.0751: 2, L0758: 2,	S0242: 2, H0223: 1,	H0685: 1, S0116: 1,	S0282: 1, H0662: 1,	S6022: 1, H0574: 1,	H0486: 1, H0013: 1,	H0098: 1, H0575: 1,	H0581: 1, H0052: 1,	.0471: 1, H0024: 1,	H0510: 1, H0594: 1,	H0328: 1, H0169: 1,	H0598: 1, H0163: 1,	10646: 1, L0637: 1,	0768: 1, L0794: 1,	.0804: 1, L0775: 1,	•	.0526: 1, L0783: 1,		.0647: 1, L0666: 1,	0664: 1, S0374: 1,	.0438: 1, H0519: 1,	H0670: 1, H0660: 1,	H0648: 1, S0378: 1,	H0696: 1, S0432: 1,	20777: 1, L0755: 1,	S0308: 1, L0591: 1,	.0608: 1, S0192: 1,	S0194: 1, H0543: 1,
207	104		202	HOC	202	99S	70H)OH	HO	707	HO	HO	0H)	HOC	2	80 <u>1</u>	907	<u> </u>	80 <u>1</u>	901	200	<u>1</u>	90H	HOC	HOC	LOJ	503	907	S01
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H0423: 1 and S0412: 1.		L0731: 7, S0358: 6, L0439: 6, L0769: 5,	L0809: 4, L0747: 4,	S0354: 3, H0580: 3,	H0438: 3, H0581: 3,	L0770: 3, L0766: 3,	L0774: 3, L0775: 3,	L0740: 3, L0749: 3,	L0758: 3, L0596: 3,	H0677: 3, H0392: 2,	H0457: 2, L0471: 2,	H0510: 2, H0169: 2,	H0551: 2, T0042: 2,	H0494: 2, S0438: 2,	L0761: 2, L0657: 2,	L0666: 2, L0565: 2,	L0438: 2, H0521: 2,	L0748: 2, L0750: 2,	L0779: 2, L0752: 2,	L0759: 2, L0588: 2,	L0605: 2, L0592: 2,	L0581: 2, L0615: 1,	S0116: 1, H0638: 1,	S0418: 1, S0444: 1,	S0360: 1, S0222: 1,	H0586: 1, H0427: 1,	H0618: 1, H0390: 1,
		Leu-9 to Val-14, Glu-41 to Ala-49.																									
	3554	3555																		,-							
	39 - 185	2-217											-										·				:
	1379	i														 											
	HOCPG48R	HOCPH40R	-																								
	HOCPG48	НОСРН40														 			-								

H0318: 1, H0007: 1, H0052: 1, H0544: 1, H0024: 1, H0014: 1, H0083: 1, H0375: 1, S6028: 1, H0266: 1, H0135: 1, H0087: 1, S0344: 1, S0426: 1, H0529: 1, L0763: 1, L0764: 1, L0662: 1, L0768: 1, L0662: 1, L0659: 1, L0518: 1, L0659: 1, L0518: 1, H06519: 1, H0547: 1, H0690: 1, H0682: 1, H0660: 1, S0328: 1, H0660: 1, S0328: 1, H0645: 1, L0777: 1, H0445: 1, L0362: 1,					
	Gly-18 to Asp-23, Gln-34 to Gly-40, Asp-81 to Lys-87.		Asp-43 to Gly-48.		
•	3556	3557	3558	3559	3560
	3 - 281	3 - 245	2 - 232	285 - 530	2 - 136
	1381	1382		1384	1385
	HOCPH44R	HOCP128R	HOCPI53R	HOCPL59 HOCPL59R	HOCPM39R
	НОСРН44	HOCP128	HOCPI53	HOCPL59	HOCPM39

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L0747: 10, L0805: 6,	5: 4,	19: 4,	40: 3,	76: 3,	50: 2,	71: 2,	69: 2,	L0770: 2,	L0761: 2,	L0794: 2,	06: 2,	L0809: 2,	547: 2,	560: 2,	521: 2,	48: 2,	58: 2,	02: 1,	657: 1,	120: 1,	42: 1,	574: 1,	486: 1,	114: 1,	318: 1,	051: 1,	1292: 1,	031: 1,	163: 1,
: 10, L(4, 1.077	,0655: 4, L0749: 4,	H0656: 3, S0440: 3,	,0803: 3, L0776: 3,	,0789: 3, H0650: 2	H0662: 2, L0471: 2	10620: 2, H0169: 2,	2, L07	2, L07	2, 1.07	.0804: 2, L0806:	2, L08	0665: 2, H0547: 2,	S0126: 2, H0660: 2,	H0648: 2, H0521: 2,	S0406: 2, L0748: 2,	.0750: 2, L0758: 2,	.0581: 2, T0002: 1	H0686: 1, H0657:	S0418: 1, S0420: 1	S0356: 1, S0442:	S0444: 1, H0574: 1,	H0559: 1, H0486: 1,	.0586: 1, T0114:	H0250: 1, H0318:	H0014: 1, H0051:	H0266: 1, H0292:	H0428: 1, H0031:	H0673: 1, H0163:
L0747	S0358: 4, L0775: 4,	L0655:	H0656:	L0803:	L0789:	H0662:	H0620:	H0616: 2,	L0769: 2.	L0800: 2,	L0804:	L0657: 2,	T0665:	S0126:	H0648	S0406	L0750	L0581	H0686	S0418	S0356	S0444	H0555	L0586	H025(H0014	H026	H0428	H067
o Ala-9																													
Ala-2 to Ala-9.																													<u>.</u>
3561				:	:																								
3 - 275				:																									
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H0591: 1, H0494: 1, L0640: 1, L0763: 1, L0772: 1, L0773: 1, L0774: 1, L0773: 1, L0384: 1, L0793: 1, H0144: 1, H0703: 1, S0374: 1, H0593: 1, H0658: 1, S0328: 1, S0152: 1, L0744: 1, L0779: 1, L0731: 1, L0759: 1, S0026: 1, H0136: 1, H0542: 1 and H0422: 1,			L0754: 4, H0290: 2, L0761: 2, L0794: 2, L0766: 2, L0793: 2, L0731: 2, L0758: 2, L0759: 2, L0594: 2, H0713: 1, H0402: 1, S0418: 1, S0354: 1, H0580: 1, L0717: 1,
	Glu-16 to Lys-21, Asn-62 to Arg-68, Asp-94 to Thr-102, Gly-161 to Leu-170.		Ser-9 to Glu-16, Phe-49 to Tyr-58, Asn-102 to Gly-110.
	3562	3563	3564
	3 - 587	26 - 358	2 - 394
	1387	1388	1389
	HOCPP80R	HOCPQ24R	носрозяк.
	HOCPP80	_	

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H0643: 1, H0574: 1, S0280: 1, H0618: 1, H0052: 1, H0546: 1, H0545: 1, H0014: 1, L0521: 1, L0773: 1, L0521: 1, L0803: 1, L0788: 1, H0670: 1, H0660: 1, H0521: 1, S0404: 1, S0406: 1, H0436: 1, L0779: 1,	HUS43: 1 and HUSSZ: 1.					L0748: 9, H0556: 8,	L0776: 4, H0547: 4,	H0521: 4, L0750: 4,	L0752: 4, L0758: 4,	S0434: 4, H0542: 4,	H0657: 3, S0418: 3,	H0586: 3, H0587: 3,	H0013: 3, H0052: 3,
	Asp-24 to Leu-31, Ile-123 to Gln-128.		Ser-11 to Trp-18,	Glu-43 to Leu-50, Phe-99 to Arg-120, Ser-136 to Tyr-144,	Glu-154 to Cys-163.	Gln-14 to Lys-19,							
	3565	3566	3567			3568							
	2 - 427	2 - 487	3 - 536			1 - 588							
	1390	1391	1392			1393							
	HOCPQ66R	HOCPO75R	HOCPR01R			HOCPR29R							
	НОСРQ66	HOCPO75				HOCPR29					-		

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H0038: 3,	10494: 3,	.0774: 3,	.0659: 3,	.0744: 3,	,0755: 3,	10543: 3,	0476: 2,	10194: 2,	10014: 2,	30003: 2,	10135: 2,	0142: 2,	.0662: 2,	.0381: 2,	.0666: 2,	10658: 2,	30044: 2,	0751: 2,	L0596: 2,	0591: 2,	.0595: 2,	10423: 2,	0134: 1,	0661: 1,	10125: 1,	0356: 1,	944: 1,	0619: 1,	10549: 1,
H0083: 3, H0038: 3,	10634: 3, I	S0440: 3, L0774: 3,	.0775: 3, L0659: 3,	10522: 3, L0744: 3,	0749: 3, I	L0759: 3, H0543: 3,	0358: 2, S	H0156: 2, H0194: 2,	.0471: 2, H0014: 2,	10201: 2, S0003: 2,	F0023: 2, H0135: 2,	H0551: 2, S0142:	.0372: 2, L0662: 2,	.0767: 2, L0381: 2,	.0383: 2, L.0666: 2,	.0665: 2, H0658: 2,	H0660: 2, S0044: 2,	S3014: 2, L0751: 2,	.0754: 2, L	.0588: 2, L0591: 2,	.0592: 2, L0595: 2,	.0601: 2, H0423: 2,	S0040: 1, S0134: 1,	S0298: 1, H0661: 1	H0664: 1, H0125: 1,	S0420: 1, S0356: 1,	S0354: 1, S0444: 1,	S0360: 1, H0619: 1,	H0393: 1, H0549: 1
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0222: 1,	0431: 1,	10574: 1,	0114: 1,	10427: 1,	10004: 1,	10581: 1,	10596: 1,	[0009: 1,	10050: 1,	0010: 1,	10288: 1,	H0039: 1,	,0055: 1,	10090: 1,	10040: 1,	10413: 1,	0438: 1,	30210: 1,	10529: 1,	.0762: 1,	.0770: 1,	L0764: 1,	L0386: 1,	.0524: 1,	.0379: 1,	.0657: 1,	.0809: 1,	.0663: 1,	30374: 1,
H0550: 1, S0222: 1	S6014: 1, H0431: 1,	H0592: 1, F	T0039: 1, T	H0075: 1, H0427: 1	H0590: 1, F	H0318: 1, H0581: 1	H0251: 1, H0596: 1,	T0115: 1, H	H0081: 1, F	H0024: 1, T	H0687: 1, H0288:	H0290: 1, H0039:	H0622: 1, L0055: 1	H0032: 1, H0090: 1	H0591: 1, F	H0063: 1, H0413: 1	T0069: 1, S0438:	H0538: 1, S0210:	S0426: 1, H0529: 1	L0520: 1, I	L0763: 1, L0770:	L0374: 1, I	L0768: 1, I	L0803: 1, I	L0806: 1, L0379:	L0527: 1, L0657: 1	L0382: 1, L0809: 1	L0788: 1, I	S0052: 1, S0374:
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H0691: 1, H0682: 1, H0435: 1, H0659: 1, S0328: 1, S0330: 1, S0152: 1, H0436: 1, L0611: 1, S0027: 1, S0028: 1, L0747: 1, L0779: 1, L0780: 1, L0731: 1, L0608: 1, S0026: 1, H0653: 1, S0192: 1, S0194: 1, H0422: 1 and S0384: 1				L0766: 2, H0660: 2,	L0748: 2, L0740: 2,	H0328: 1, L0769: 1,	L0651: 1, L0653: 1,	H0593: 1 and S0126: 1.										H0620: 2, S0022: 2,
		Glu-22 to Glu-30,	Lys-42 to Glu-51.	Pro-7 to Ser-36,	Cys-49 to Pro-57,	Pro-85 to Ala-91,	Gly-97 to Pro-109.				Tyr-1 to Gly-39,	Ala-41 to Tyr-67.			Ser-49 to Asp-56;	Arg-64 to Ile-71,	Pro-80 to Ile-89.	His-10 to Asp-20,
	3569	3570		3571					3572	3573	3574		3575	3576	3577			3578
	168 - 335	62 - 256		3 - 440					1 - 279	1 - 102	3 - 203		1 - 225	333 - 560	3 - 269			339 - 599
	1394	1395		1396					1397	1398	1399		1400	1401	1402			1403
	HOCPR33R	HOCPR53R		HOCPR55R					HOCPR77R	_ 4	HOCPS35R		HOCPU03R	HOCPU30R	HOCPU68R			HOCPV29R
	HOCPR33	HOCPR53		HOCPR55					HOCPR77	HOCPR82	HOCPS35		HOCPU03		HOCPU68			HOCPV29

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L0438: 2, T0002: 1,	H0050: 1, H0373: 1,	H0051: 1, H0028: 1,	S0036: 1, H0551: 1,	H0059: 1, S0150: 1,	L0662: 1, S0126: 1,	H0435: 1, H0660: 1,	S0044: 1 and H0694: 1.	L0803: 7, L0439: 5,	L0809: 4, L0749: 4,	L0776: 3, H0520: 3,	L0748: 3, L0750: 3,	H0543: 3, S0440: 2,	L0794: 2, L0804: 2,	L0438: 2, H0547: 2,	H0660: 2, S0406: 2,	L0743: 2, L0777: 2,	L0759: 2, H0556: 1,	H0656: 1, S0358: 1,	S0360: 1, H0637: 1,	H0580: 1, H0741: 1,	H0733: 1, H0619: 1,	H0415: 1, H0600: 1,	H0587: 1, H0486: 1,	H0013: 1, H0427: 1,	S0474: 1, H0581: 1,	H0263: 1, H0457: 1,	H0615: 1, H0428: 1,	H0644: 1, H0032: 1,
Asn-27 to Gly-33.			-										•															
	_							3579																				
					-			1 - 390	1			_																
						···-		1404																				
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S0036: 1, H0591: 1, H0634: 1, H0264: 1, S0438: 1, S0422: 1, S0002: 1, L0520: 1, L0766: 1, L0649: 1, L0775: 1, L0655: 1, L0775: 1, L0655: 1, L0790: 1, L0666: 1, H06593: 1, H0682: 1, H0670: 1, H0631: 1, S0013: 1, H0631: 1, S0206: 1, L0756: 1, L0731: 1, L0608: 1,	
	lle-53 to Trp-63, Ala-80 to Gly-89, Arg-134 to Gln-139.
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S0420: 1, S0376: 1,	S0360: 1, S0007: 1,	S0300: 1, S0222: 1,	S6014: 1, H0250: 1,	L0021: 1, L0022: 1,	S0049: 1, H0309: 1,	H0150: 1, H0009: 1,	S0051: 1, H0188: 1,	F0006: 1, H0604: 1,	H0673: 1, H0169: 1,	H0087: 1, L0065: 1,	L0763: 1, L0638: 1,	L0773: 1, L0662: 1,	L0768: 1, L0766: 1,	L0649: 1, L0651: 1,	L0657: 1, L0540: 1,		L0530: 1, L0787: 1,	L0789: 1, L0665: 1,	S0374: 1, S0126: 1,	H0684: 1, H0660: 1,	H0539: 1, S0380: 1,	H0576: 1, L0741: 1,	L0742: 1, L0749: 1,	L0750: 1, L0779: 1,	L0759: 1, S0031: 1 and	L0590: 1.			S0358: 45, S0360: 30,
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				Met-1 to Leu-6,	Gly-19 to Lys-26,	Ala-32 to Lys-38,	Ala-89 to Leu-94,	Gly-134 to Val-141.	Met-36 to Pro-42,	Pro-49 to Thr-55,	Leu-104 to Met-110,	Trp-112 to Asn-117,	Glu-122 to Tyr-134,	Gln-142 to Ser-148,	Leu-174 to Tyr-180.
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S0116: 1, S0298: 1,	S0410: 1, S0468: 1,	H0369: 1, H0453: 1,	370: 1, H0392: 1,	H0602: 1, H0587: 1,	70039: 1, H0013: 1,	H0004: 1, H0052: 1,	309; 1, H0046; 1,	H0041: 1, H0081: 1,	H0011: 1, H0049: 1,	[10079: 1, H0375: 1,	250: 1, S0214: 1,	H0252: 1, H0428: 1,	0006: 1, H0032: 1,	H0708: 1, H0316: 1,	0598: 1, H0400: 1,	H0100: 1, H0494: 1,)475: 1, S0014: 1,	S0015: 1, H0625: 1,	S0382: 1, H0646: 1,	.0762: 1, L0763: 1,	.0630: 1, L0646: 1,	.0768: 1, L0649: 1,	_	L0809: 1, L0647: 1,	.0663: 1, L0665: 1,	S0374: 1, H0691: 1,	H0547: 1, H0682: 1,	H0521: 1, S0027: 1,
OS	OS SO	OH H	OH HO	OH H		OH)	OH)HI)[1]	TO	os	H		H	H	H	그)S	SC	<u>급</u>	<u> </u>					Š	32 :	H .
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L0748: 1, L0756: 1, L0777: 1, L0752: 1, L0755: 1, H0343: 1, H0707: 1, L0605: 1, H0668: 1, S0196: 1 and	H0046: 12, L0771: 7, L0806: 3, L0666: 3, L0866: 3, L0666: 3, L0565: 3, L0748: 3, S0444: 2, S0408: 2, H0393: 2, L0372: 2, L0662: 2, L0766: 2, L0744: 2, L0731: 2, H0294: 1, S0442: 1, S0278: 1, H0549: 1, H0297: 1, H0549: 1, H0252: 1, H0383: 1, H0252: 1, H0100: 1, H0494: 1, H0413: 1, L0772: 1, L0761: 1, L0772: 1, L0764: 1, L0800: 1, L0764: 1,
	Leu-16 to Pro-22, Val-30 to Asp-36.
	3604
	61 - 546
	1429
	HOCQJ35R
	носолз

				·
L0649: 1, L0803: 1, L0805: 1, L0653: 1, L0515: 1, L0367: 1, L0665: 1, S0374: 1, H0684: 1, H0660: 1, S0328: 1, S0044: 1, S0404: 1, L0740: 1, L0747: 1, L0757: 1, L0758: 1, S0260: 1 and S0276: 1.			H0328: 1 and H0435: 1.	L0493: 4, L0794: 3, H0497: 2, H0038: 2, L0766: 2, L0809: 2, L0779: 2, L0777: 2, L0759: 2, L0601: 2, H0170: 1, S0114: 1, H0341: 1, H0125: 1, S0046: 1, H0592: 1, H0587: 1, H0559: 1, H0587: 1, H0529: 1, H0538: 1, H0386: 1, H0538: 1, H0529: 1,
	Gly-2 to Gly-12, Gly-14 to Asn-28, Met-32 to Gly-49, Lys-54 to Phe-76.		Ala-1 to Ser-8.	Gln-1 to Pro-16.
	3605	3606	3607	3608
	3 - 230	2-76	234 - 395	266 - 484
	1430	1431	1432	1433
	HOCQM24 HOCQM24R	HODAF78R		HODCT68R
	HOCQM24	HODA F78	HODBT55	норст68

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L0775: 1, L0666: 1, H0659: 1, H0518: 1, L0741: 1, L0439: 1, L0747: 1, L0593: 1, L0594: 1, H0665: 1 and H0667: 1.	AR089: 19, AR104: 18, AR060: 12, AR096: 7, AR055: 4, AR061: 3 L0751: 29, H0549: 17, H0550: 17, L0666: 9, L0747: 8, H0682: 7, H0640: 7, H0682: 7, H0640: 7, H0682: 7, H0650: 4, S0330: 4, S0044: 4, L0362: 4, H0597: 3, H0688: 3, L0764: 3, L0753: 3, L0758: 3, L0753: 3, L0758: 3, L0753: 3, H0431: 2, H0046: 2, H0647: 2, L0774: 2, L0805: 2, L0774: 2, L0805: 2, L0778: 2, L0805: 2, L0778: 2, H0648: 2, H0672: 2, S0328: 2, S0404: 2, L0744: 2, L0779: 2,
	3609
	2-139
	1434
·	HODCZ52R
·	HODCZ52

S0436: 2, H0662: 1, S0442: 1, S0444: 1, S0408: 1, S0408: 1, H0045: 1, H00609: 1, L06009: 1, L06009: 1, L06009: 1, L06009: 1, L06009: 1, L0663: 1, L06	L0794: 4, L0803: 4, H0328: 3, L0779: 3, L0515: 2, L0809: 2, H0144: 2, L0754: 2, L0756: 2, L0759: 2, H0171: 1, S0282: 1, H0346: 1, S0222: 1, H069: 1, H0486: 1, H069: 1, H056: 1, L0598: 1, L0638: 1, L0766: 1, L0650: 1,
	Gly-7 to Glu-41.
	3610
	3 - 194
	1435
	HODDIS7R
	HODDIS7

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D	
L0774: 1, L0805: 1, L0776: 1, L0657: 1, L0559: 1, L0659: 1, L0543: 1, L0438: 1, H0682: 1, S0328: 1, L0745: 1, L0750: 1, L0777: 1, L0752: 1, L0778: 1, H0343: 1 and H0665: 1.	L0754: 20, L0759: 6, L0794: 4, H0615: 3, L0438: 3, S0126: 3, S0360: 2, H0411: 2, H0050: 2, L0748: 2, L0749: 2, L0749: 2, L0749: 2, L0749: 2, L0749: 2, L0749: 2, L0755: 2, H0595: 2, L0755: 2, H0595: 2, L0361: 2, H0638: 1, S0376: 1, H0628: 1, H0625: 1, H0625: 1, L0662: 1, L0764: 1, L0662: 1, L0669: 1, L0669
·	Leu-7 to Thr-16.
÷	3611
	11 - 295
	1436
·	HODEJ47R
	HODEJ47

1 00	830: 1, 530: 1, 731: 1, L0605: 1.		0439: 4, 749: 3,	769: 2,	774: 2,	777: 2,	142: 1,	767: 1,	784: 1,	789: 1,	438: 1,	i L0589: 1.	nd S0031: 1.			10615: 1,	791: 1,	0752: 1 and				nd H0615:	
30.1	L0789: 1, L0663: 1, H0144: 1, S0330: 1, S0028: 1, L0751: 1, L0750: 1, L0731: 1, L0758: 1 and L0605: 1		L0803: 7, L0439: 4, L0663: 3, L0749: 3,	H0615: 2, L0769: 2,	L0794: 2, L0774: 2,	L0748: 2, LO	L0752: 2, L0	H0646: 1, L0	L0804: 1, LO	L0805: 1, L0789: 1,	L0666: 1, L0	H0670: 1 and L0589:	H0615: 1 and S0031:			L0662: 2, H0615: 1,	L0761: 1, L0791: 1,	H0519: 1, L(L0758: 1.	H0615: 2		H0328: 1 and H0615:	1.
													Asn-1 to Ser-6,	Thr-22 to Asn-32.						Ser-1 to Ser-9, Asn-15 to Ala-21.			
		3612	3613										3614		3615	3616				3617	3618	3619	
		1 - 54	187 - 432										349 - 444		92 - 289	2 - 421				153 - 347	370 - 546	1	
		1437	1438				-	-					1439		1440	1441	•			1442	1443	4	
		HODEO42R	1				-	_					HODFO42R	ייי איז איז איז איז איז איז איז איז איז	HODFR57R	HODESCOR	NOOT TOLL			HODEU47R	TIONEVICOD	HODEV89 HODEVSAR 1444	
		CNOTOTI											UODEO42	מבאממטם	TODED 57	TODESEO	HODESOO			HODEU47	O) Indiana	HODEV69	

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	H0615: 2 and L0532: 1.	H0615: 4 and L0659: 1.		L0747: 16, L0754: 10,	L0751: 7, L0757: 7,	H0657: 6, L0748: 6,	L0771: 5, H0659: 5,	L0740: 5, L0755: 5,	H0594: 4, L0744: 4,	H0341: 3, S0358: 3,	S0278: 3, H0333: 3,	H0615: 3, S0366: 3,	L0520: 3, L0775: 3,	L0517: 3, S0374: 3,	S0328: 3, L0731: 3,	H0136: 3, T0002: 2,	S0218: 2, S0116: 2,	S0354: 2, S0376: 2,	H0637: 2, S0045: 2,	H0370: 2, L0622: 2,	H0156: 2, H0599: 2,	H0545: 2, H0266: 2,	H0039: 2, H0617: 2,	H0132: 2, L0769: 2,	L0665: 2, H0672: 2,	H0539: 2, H0134: 2,	L0756: 2, L0752: 2,
	-	<i>:</i>		Ala-9 to Asp-21,	Thr-31 to Arg-37.																						
3620	3621	3622	3623	3624																						:	
309 - 566	220 - 423	58 - 231	1 - 432	1 - 207			•													-		-				:	
1445	1446	1447	1448	1449													-										
HODEW79 HODEW79R	HODEX45 HODEX45R	HODEX69R	HODEX79R	HODFC44 HODFC44RP	8																		-	-			
HODEW79	HODEX45	норех69	HODEX79	HODFC44																							

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.0758: 2, H0624: 1,	H0170: 1, H0265: 1,	S6024: 1, H0656: 1,	H0255: 1, H0661: 1,	H0125: 1, H0192: 1,	8: 1, S0420: 1,	30360: 1, H0580: 1,	93: 1, S0300: 1,	H0411: 1, H0369: 1,	61: 1, S0222: 1,	.0623: 1, S0280: 1,	40618: 1, H0318: 1,	H0581: 1, H0421: 1,	.0738: 1, H0086: 1,	_0471: 1, H0024: 1,	82: 1, L0043: 1,	S0214: 1, H0031: 1,	H0673: 1, H0068: 1,	138: 1, H0634: 1,	H0087: 1, T0067: 1,	H0412: 1, H0100: 1,	H0512: 1, S0144: 1,	S0142: 1, S0344: 1,	S0422: 1, H0517: 1,	L0762: 1, L0763: 1,	(61: 1, L0764: 1,	768: 1, L0766: 1,	L0649: 1, L0774: 1,	.0651: 1, L0776: 1,	.0655: 1, L0526: 1,
1075	H017	8602	H025	H012	S041	903	H03	H04]	H02(7907	90H	H05	1.07	1.047	00H	802	90H	00H	00H	H04	H05	201	804	107	107	107	907	<u>100</u>	100
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L0518: 1, L0783: 1, L0809: 1, L0666: 1, L0663: 1, H0144: 1, H0593: 1, H0682: 1, H0648: 1, S0044: 1, H0214: 1, L0439: 1, L0749: 1, L0750: 1, L0753: 1, H0444: 1, H0445: 1, L0593: 1 and H0542: 1.	L0794: 3, L0776: 2, S0358: 1, H0615: 1, H0538: 1, L0662: 1, L0803: 1, L0805: 1, L0791: 1, L0663: 1, S0374: 1, H0658: 1 and L0758: 1.		L0766: 2, H0328: 1 and H0615: 1.		H0328: 1 and H0615: 1.			H0615: 2, H0370: 1, H0328: 1 and L0749: 1.
	Ser-22 to Gln-38.			3628 Phe-45 to Tyr-60, Arg-67 to Glu-74.	Pro-18 to Pro-23, Pro-37 to Val-42.		Lys-69 to Asp-81.	•
	3625	3626	3627	3628	3629	3630	3631	3632
	3 - 281	296 - 433	293 - 409	2 - 259	97 - 225	1 - 69	16 - 324	1 - 165
·	1450	1451	1452	1453	1454	1455		1457
	норгр90к	HODFE81R	HODFI68R	HODF114R	HODFJ52R	HODFK30 HODFK30RA	HODFK89R	HODFL50R
	норғр90	HODFE81	норя	HODFJ14	HODF152	HODFK30	HODFK89	HODFL50

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	H0615: 2	L0666: 3, H0615: 2, S0116: 1, H0675: 1, H0012: 1, L0055: 1,	S0438: 1, L0763: 1, L0667: 1, L0649: 1, I 0803: 1, L0804: 1,	L0659: 1, H0689: 1, H0659: 1, L0752: 1 and S0026: 1.								H0615: 2			1 11000	H0328: 1, H0615: 1 and L0365: 1.	H0615: 3, H0693: 2,	
Ala-2 to Trp-9.		Glu-12 to Gly-27, Gln-37 to Tyr-49.	<u> </u>		His-33 to Leu-39, Gly-49 to Glu-56,	Ser-60 to Lys-143.	Leu-33 to Val-40, Lys-144 to Glu-152.	His-1 to Lys-12, Gln-17 to Leu-27,	Val-69 to Asp-76,	Asp-122 to Trp-128,	Ser-155 to His-164.				Pro-1 to Asp-7.			
3633	1-	1			3636		3637	3638				3639	3640	3641	3642	3643	2644	<u>-</u>
204 - 303	251 - 565	1 - 195			11 - 439		1-519	2 - 535				84 - 512	<u> </u>	2 - 130	291 - 467	7 - 168	010	
1450	1459	1460			1461		1462	1463				1464	4—					1469
	+	1			HODFP51R		норго19 норго19к	HODFQ37R				HODER72R	-	HODFU78R	HODFV22R	HODFV60R		HODFV69K
	HODFL/S HODFL/SR	HODFO64			HODFP51		НОДРО19	HODFQ37				HODEP 72	HODE 194	HODE 178	HODEV22	HODFV60		HODFV69

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H0266: 1, S0002: 1, L0790: 1, H0547: 1, S0152: 1, H0521: 1, H0696: 1, L0366: 1, S0194: 1 and S0276: 1.				H0615: 2, S0444: 1,	L0438: 1 and L0439: 1.						,			H0615: 2	L0754: 2, L0756: 2,	H0615: 1, L0438: 1 and	L0747: 1.	L0741: 4, H0615: 2,	L0769: 2, L0793: 2,	H0328: 1, S0144: 1,	L0794: 1, L0499: 1,	L0805: 1, L0438: 1 and	L0755: 1.
				Lys-8 to Tyr-26,	Thr-67 to Met-73.	Val-1 to Ser-9,	Gln-22 to Ser-29.		Thr-14 to Asn-19.	,	Thr-7 to Ser-14,	Pro-28 to Gly-36.	His-8 to Gly-18, Phe-25 to Gly-30.					Lys-1 to Thr-7,	Ser-13 to Arg-33,		Phe-143 to Thr-148.		
	3645	3646	3647	3648		3649		3650	3651	3652	3653		3654	3655	3656			3657				v	
	360 - 623	14 - 154	587 - 718	22 - 309		186 - 308		3 - 95	91 - 225	129 - 407	25 - 132		36 - 128	2 - 166	127 - 300	•		2 - 583					
	1470	1471	1472	1473		1474		1475	1476	1477	1478		1479	1480	1481			1482				,	
	HODFV79R	HODFV82R	HODFV95R	HODFY38R		HODGA29R		HODGB69R	HODGB78R	HODGE22R	HODGH30 HODGH30R		HODGH43R	HODGH65R	HODGL65R			HODGN53R					
	HODFV79	HODFV82	HODFV95	HODFY38		HODGA29		HODGB69	HODGB78	HODGE22	норен30		нордн43 нордн43	НОДСН65	HODGL65			HODGN53					

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	H0615: 2 and L0779:								H0615: 3 and L0766:	1.			L0805: 19, L0794: 13,	L0766: 12, L0740: 11,	L0747: 11, L0779: 11,	L0803: 8, H0521: 7,	L0439: 7, L0731: 7,	L0605: 7, S0474: 6,	S0422: 6, L0769: 6,	L0776: 5, L0665: 5,	L0748: 5, L0758: 5,	H0618: 4, S0010: 4,	L0771: 4, L0789: 4,	S3014: 4, H0341: 3,	H0615: 3, L0659: 3,	S0052: 3, L0438: 3,
		Glu-23 to Val-28,	Arg-56 to Ser-64.				Lys-1 to Gly-12, Pro-25 to Thr-37.	Ser-46 to Gly-51.					Pro-30 to Asn-35,	Arg-39 to Asn-57,	Ser-62 to Ile-67,	Glu-73 to Arg-81,										
3658	3659	3660		3661	3662	3663	3664		3665		3666	3667	3668													
361 - 561	1 - 348	2 - 193		1 - 198	9 - 131	205 - 354	182 - 3		358 - 513		219 - 572	379 - 573	1 - 504													
1483	1484	1485	•	1486	1487	1488	1489	•	1490		1491	1492	1493													
HODGO02R	HODGO09 HODGO09R	HODGO21 HODGO21R		HODGP53R	HODGQ52 HODGQ52R	HODGS83R	HODGW91 HODGW91R		HODGX46 HODGX46R		HODGX91R	HODGY90 HODGY90R	HODGZ63R													
HODGO02	НОДСО09	HODG021		HODGP53	HODGQ52	HODGS83	HODGW91		HODGX46		HODGX91	HODGY90	HODGZ63												,	

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043	0759	3222	<u> 1059</u>	1005	020	1003	080	037	0682	470	L0749: 2,	0424	,000	.078	4	l, H0733:	043	028	90	H0251:	1012	1063	S0003:	.0483:	460	003	1004	1041	8
3, H	3, L	2, S(2, H	2, H	2, H	2, H	2, L	2, S	2, H	2, L	2, L	2,8	1, T	1, L	1, S	1, H	1, H	1, S	1, S	1, H	1, H	1, H	1, S		1, H	1, S	1, E	1,	1,5
H0658: 3, H0436: 3,	.0752: 3, L0759: 3,	S0360: 2, S0222: 2,	H0574: 2, H0599: 2	H0196: 2, H0050: 2,	H0266: 2, H0708: 2,	H0135: 2, H0038: 2,	H0616: 2, L0804: 2,	H0144: 2, S0374: 2,	.0352: 2, H0682: 2,	.0743: 2, L0744: 2,	.0751: 2,]	.0756: 2, S0424: 2	H0624: 1, T0002:	H0657: 1, L0785:	H0671: 1, S0442:	S0408: 1	S0278: 1, H0431: 1	H0013: 1, S0280: 1	581:	H0052: 1,	,0040: 1, H0123:	.0471: 1, H0630:	H0375: 1,	H0252: 1,]	F0006: 1, H0644:	H0673: 1, S0036: 1	H0163: 1, H0040:	H0551: 1, H0412: 1	H0413: 1, S0038:
H	<u>2</u>	803	유	유	丑	유	丑	유	9	<u>i</u>	<u>ප</u>	<u>9</u>	ឣ	H	표	80	80	유	유	ឣ	<u>ឧ</u>	<u>ន</u>	ឣ	H	은	H	Ħ	H	뵈
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H0509: 1, H0131: 1, \$80344: 1, L0598: 1, L0369: 1, L0763: 1, L0667: 1, L0662: 1, L0770: 1, L0662: 1, L075: 1, L0768: 1, L075: 1, L0375: 1, L0806: 1, L0653: 1, L0655: 1, L0809: 1, \$80126: 1, H0689: 1, H0684: 1, H0660: 1, H0672: 1, S0146: 1, \$3012: 1, S0027: 1, \$3012: 1, S0027: 1, \$1, L0745: 1, L0750: 1, L0745: 1, L0750: 1, L0745: 1, L0750: 1, L0361: 1, H0653: 1, H0442: 1, H0653: 1, H0423: 1 and H0506: 1.			H0615: 2	H0328; 1 and H0615; 1.	:	
: : :		Pro-8 to Asn-18, Gln-25 to Glu-30.	Tyr-33 to Tyr-38.	•	Glu-7 to Tyr-12, Glu-44 to Asp-49, Pro-57 to Gly-64,	Leu-69 to Arg-74,
	3669	3670	3671	3672	3673	
	12 - 179	3 - 134	14 - 205	190 - 366	3 - 515	•
	1494	1495	1496		1498	
	HODGZ76R	+	HODHD64R	норнезе норнезек	HODHES4 HODHES4R	
	92250cn	HODHD23	DAUDIOU LA	HODHE36	HODHE54	

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	H0328: 1 and H0615: 1.										H0261: 1 and H0615:	•	AR104: 14, AR096:	13, AR089: 11, AR060:	10, AR055: 6, AR061:		H0615: 3, H0658: 2,	HUGOI: 1 and LU333: 1.	,						S0360: 9, L0642: 5,	L0752: 5, L0662: 4,	L0659: 4, L0666: 4,
Thr-76 to Tyr-86.	Lys-42 to Asp-48.	Glu-11 to Trp-17,	Val-22 to Ser-29,	Asp-45 to Lys-52.	Trp-12 to Ile-18,	Gln-40 to Leu-54.	Lys-17 to Ala-24.	Glu-1 to Gln-7,	Met-64 to Gly-69.		Lys-9 to Ser-15,	Val-42 to Thr-50.	Phe-116 to Met-126. AR104: 14, AR096:					4		Gln-40 to Glu-48,	Phe-68 to Arg-77,	Ile-108 to Glu-113.		Arg-33 to Thr-38.	Г	.	
	3674	3675			3676		3677	3678		3679	3680		3681				,		3682	3683			3684	3685	3686		
	2 - 166	82 - 300			281 - 499		71 - 238	2 - 292		2 - 115	218 - 472		31 - 594						504 - 746	3 - 344			73 - 216	1 - 159	1 - 486		
	1499	1500			1501		1502	1503		1504	1505		1506						1507	1508			1509	1510	1511		
	HODHE88R	норндзе норндзек			HODHJ56R	_	HODHK82 HODHK82R	норнк86 норнк86к		HODIG29R	≴		HODJZ09R			-			HODKB19R	HODKB82R			HODKB89R	HODKB93R	HODKC29R		
	норне88	HODHG56			норн156		НОДНК82	норнк86		HODIG29	норлгзе норлгзе		HODIZ09						HODKB19 HODKB1	HODKB82			-	HODKB93	HODKC29		

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S0007: 3, L0646: 3, L0776: 3, L0750: 3, H0662: 2, S0444: 2, S0222: 2, L0483: 2,	L0764: 2, L0046: 2, L0775: 2, L0665: 2, H0648: 2, L0779: 2, S0031: 2, S0110: 1, H0638: 1, S0418: 1,	S0376: 1, S0408: 1, L0717: 1, H0331: 1, H0574: 1, S0414: 1, H0497: 1, H0156: 1.	L0021: 1, S0049: 1, H0310: 1, H0327: 1, H0373: 1, T0010: 1,	S6028: 1, H0615: 1, S0366: 1, S0036: 1, H0038: 1, H0616: 1,	H0413: 1, T0041: 1, H0494: 1, S0438: 1,	S0440: 1, H0509: 1, S0142: 1, S0210: 1,	L0598: 1, L0762: 1, L0640: 1, L0631: 1,	L0772: 1, L0766: 1, L0551: 1, L0774: 1,	 S0378: 1, S0152: 1,
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H0696: 1, S3014: 1, L0439: 1, L0740: 1, L0754: 1, L0756: 1, L0780: 1, L0755: 1, L0758: 1, S0434: 1 and H0667: 1.	H0615: 2		H0615: 2 and H0696: 1.	H0615: 2 and H0328: 1.			H0615: 2					•								
	A == 11 +2 Sa= 17	Arg-11 to Ser-17, Leu-51 to Pro-57.	Pro-5 to Asn-10.	Ser-18 to Ile-23.	Lys-12 to Ser-18.			Asp-40 to Gly-45,	Leu-59 to Glu-6/,	Lys-95 to Ile-105,	Asp-113 to Gly-118.	Ser-15 to Phe-23,	Lys-41 to Tyr-46,	Pro-48 to Val-59,	Thr-90 to Ile-103,	Ser-110 to Arg-116,	Pro-123 to Lys-137,	Lys-157 to Glu-167,	Arg-178 to Arg-194.	Phe-106 to Pro-114.
	3687	3088	3689	3690	3691	3692	3693	3694		٠		3695								3696
	242 - 373	43 - 213	413 - 237	314 - 412	244 - 417	28 - 240	452 - 595	2 - 415				1 - 642								3 - 344
	1512	1513	1514	1515	1516	1517	1518	1519				1520								1521
	HODKC67R	HODKD20 HODKD20K	НОДКД64 НОДКД64К	HODKE85 HODKE85R	HODKF91R	HODKG09R	HODKJ77 HODKJ77R	норкк26 норкк26к				HODKK40R								HODKK73 HODKK73R
	HODKC67	HODKD20	НОДКД64	HODKE85	HODKF91	HODKG09	HODKJ77	HODKK26				HODKK40								HODKK73

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H0615: 2				110250: 4 1 0770: 3	H0659; 4, L0770; 5, (0747; 3, H0663; 2,	S0360: 2, L0521: 2,	0776: 2, S0126: 2,	10658: 2, S0406: 2,	.0755: 2, L0758: 2,	,0759: 2, S0242: 2,	T0002: 1, H0686: 1,	H0341: 1, L0616: 1,	0354: 1, S0408: 1,	H0052: 1, H0009: 1,	H0083: 1, H0266: 1,	H0615: 1, L0483: 1,	H0040: 1, H0561: 1,	S0422: 1, L0520: 1,	L0625: 1, L0662: 1,	L0766: 1, L0649: 1,	L0519: 1, H0648: 1,	S3014: 1, L0748: 1 and	L0779: 1.				
Ser-8 to Asn-15.					<u>. </u>	S	<u></u>	<u>E</u>	<u></u>			<u> </u>												╂~~	Lys-31 to Tyr-43, 1 vs-45 to Glu-50.	Ile-96 to Lys-102,	Asp-123 to Gly-128.
3697		3699	3700	3701	3702																			3703			
272 - 433	439 - 579	36 - 149	1-210	2 - 409	20 - 328		_																	1 - 399			
1500	╁	1-	1525	1526	1527																		_	1528			
- 1-	╁		┺-	<u>_</u>	╌																			TIOE ABOTE	HOLVE		
	HODKK// HODKA//R	HODKW25 F	HODKW31 HODKW31R	HODKN65	HOECR39 HOECR39R								-					,				<u></u>			HOFAB//		

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					H0415: 2						H0415: 2						,	,										
		Arg-1 to Phe-16.	His-3 to Ala-19.		Glu-8 to Thr-14.	Ala-5 to Arg-14.	Asn-41 to Lys-56.			Asp-1 to Leu-10.	Arg-1 to Arg-8,	Ala-22 to Met-31,	Asp-51 to Gln-56.	Arg-1 to Ala-6,	Ser-68 to Arg-73,	Arg-87 to Ala-93.	Arg-1 to Cys-10,	Glu-75 to Arg-94,	Arg-99 to Gly-115.	Lys-52 to Asp-57.	Arg-1 to Phe-7,	Gly-24 to Arg-36.		Val-31 to Val-38,	Phe-46 to Ile-59,	Arg-120 to Gln-126.		Asn-49 to Leu-54.
3704	3705	3706	2018	3708	3709	3710	3711	3712	3713	3714	3715			3716			3717			3718	3719		3720	3721			3722	3723
95 - 277	85 - 219	2 - 397	1 - 162	1 - 204	69 - 323	1 - 423	50 - 289	1 - 264	1 - 369	3 - 296	2 - 481			2 - 421			25 - 387			1 - 186	2 - 232		1 - 453	2 - 682			1 - 345	3 - 326
1529	1530	1531	1532	1533	1534	1535	1536	1537	1538	1539	1540			1541			1542	_		1543	1544		1545	1546			1547	1548
HOFMF37R	HOFMF79RA	HOFMJ88R	HOFMJ93R	HOFMM84 HOFMM84R	HOFMN93R	HOFMP59R	HOFWT68R	HOFMT69R	HOFMU92R	HOFMV06R	HOFNF63R			HOFNF76R			HOFNG51R			HOFNK44R	HOFNY53R		HOFOB65R	HOFOB79RP	400		HOFOE22R	HOFOF47R
HOFMF37	HOFMF79	HOFMJ88	HOFMJ93	HOFMM84	HOFMN93	HOFMP59	HOFMT68	HOFMT69	HOFMU92	HOFMV06	HOFNF63		-	HOFNF76			HOFNG51		. ;	HOFNK44	HOFNY53		HOFOB65	HOFOB79			HOFOE22	HOFOF47

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		H0436: 44, L0748: 14, L0749: 9, L0749: 9, L0747: 9, L0769: 7, L0803: 7, L0769: 7, L0803: 7, L0766: 6, L0766: 6, L0766: 6, L0766: 6, L0779: 6, H0542: 6, L0779: 6, H0542: 6, L0779: 6, H0542: 6, L0779: 4, L0756: 3, L0779: 4, L0777: 4, L0780: 4, L0777: 4, L0780: 4, L0777: 4, L0780: 4, L0777: 4, L0780: 3, L0763: 3, H0672: 3, H0672: 3, H0673: 2, H0619: 2, H0486: 2, H0635: 2, H063
Ala-4 to Arg-11.	Ala-1 to Tyr-6, Ile-17 to Ser-32.	Pro-34 to Lys-39.
3724	3725	3726
1 - 351	35 - 193	110 - 286
15.40	1550	1551
DECEMBER	HOGAC46 HOGAC46R	HOGAQ03 HOGAQ03R
	HOFOF56 HOGAC46	HOGAQ03

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H0318: 2, S0049: 2,	H0046: 2, L0041: 2,	H0688: 2, H0428: 2,	H0628: 2, L0055: 2,	H0040: 2, H0087: 2,	H0488: 2, H0100: 2,	F0042: 2, H0529: 2,	.5566: 2, L0667: 2,	,0771: 2, L0768: 2,	, L0517: 2,	.0809: 2, L0666: 2,	H0520: 2, H0519: 2,	H0593: 2, S0380: 2,	H0694: 2, L0742: 2,	J0751: 2, L0750: 2,	S0434: 2, S0436: 2,	.0361: 2, L0601: 2,	.0603: 2, S0424: 2,	H0506: 2, H0265: 1,	H0650: 1, H0657: 1,	H0656: 1, L0760: 1,	1, S0001: 1,	H0663: 1, H0662: 1,	S0418: 1, S0356: 1,	, H0580: 1,	S0007: 1, H0229: 1,	, S0046: 1,	, H0351: 1,	H0411: 1, H0431: 1,	Н0592: 1, Н0497: 1,
H0318:	H0046:	H0688:	H0628:	H0040:	H0488:	T0042: 2	1.5566: 2	L0771: 2	1.0375: 2	T0809: 3	H0520:	H0593;	H0694:	L0751: 2	S0434: 2	L0361: 2	L0603: 2	H0506:	H0650:	H0656:	H0341:	H0663:	S0418: 1	S0408: 1	S0007: 1	\$0045: 1,	80132: 1,	H0411:	H0592:
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H0069: 1, S0280: 1, H0118: 1, H0618: 1, S0010: 1 H0421: 1	H0251: 1, H0204: 1, H0572: 1, H0050: 1,	0012: 1, H0373: 1,	H0354: 1, H0252: 1,	328: 1, H0622: 1,	H0181: 1, H0617: 1,	10673: 1, H0038: 1, 10634: 1 H0616: 1.	0155: 1, H0264: 1,	S0038: 1, H0022: 1,	H0625: 1, H0538: 1,	0210: 1, S0002: 1,	0762: 1, L0770: 1,	L5565: 1, L0373: 1,	0800: 1, L0642: 1,	0374: 1, L0764: 1,	0364: 1, L0388: 1,	L0655: 1, L0657: 1,	0526: 1, L0518: 1,	L0382: 1, L0789: 1,	0792: 1, L0664: 1,	H0723: 1, H0726: 1,	S0126: 1, H0689: 1,	10690: 1, H0682: 1,	H0435: 1, H0670: 1,	H0648: 1, S0330: 1,
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S0378: 1, S0152: 1, S0406: 1, H0555: 1, H0678: 1, L0744: 1, L0752: 1, L0753: 1, H0445: 1, H0707: 1, L0593: 1, L0595: 1, S0011: 1, H0668: 1, S0026: 1, H0665: 1, H0677: 1.	L0777: 7, L0747: 6, L0779: 6, L0748: 5, L0776: 4, L0749: 4, L0770: 3, L0766: 3, L0759: 3, S0442: 2, L0771: 2, L0761: 2, L0771: 2, L0774: 2, L0527: 2, L0783: 2, L0655: 2, L0757: 2, L0758: 2, S0412: 2, H0402: 1, S0444: 1, H0438: 1, L0471: 1, H0674: 1, L0643: 1, L0763: 1, L0643: 1, L0764: 1, L0662: 1, L0764: 1, L0662: 1, L0764: 1, L0662: 1, L0387: 1, L0677: 1, L0526: 1, L0677: 1,
	Asp-15 to His-22.
	3727
·	265 - 591
	1552
	HOGBE60R
	НОСВЕ

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H0435: 1, S0330: 1, S0406: 1, L0740: 1, L0751: 1, L0780: 1,				H0435: 4, H0726: 3, L0748: 3, H0355: 2,	H0059: 2, L0525: 2,	H0713: 1, H0717: 1,	H0586: 1, H0427: 1,	H0575: 1, H0744: 1,	H0327: 1, H0271: 1,	H0622: 1, H0087: 1,	S0438: 1, L0792: 1,	H0593: 1, H0555: 1,	L0698: 1 and H0506: 1.	H0052: 1, H0135: 1,	H0429: 1, H0519: 1 and	H0435: 1.								
THIS I												-					Val-6 to Ala-11.	Thr-56 to Glu-61,	Lys-87 to Lys-95,	Thr-110 to Arg-118.		Arg-11 to Arg-16,	Pro-46 to Thr-52,	His-58 to Pro-65,
	3728	3729	3730	3731										3732			3733	3734			3735	3736		
	230 - 436	50 - 409	58 - 207	3 - 332										10 - 480	2		3 - 254	7 - 507			3 - 143	1 - 486		
	1553	1554	1555	1556										1557	, , , ,		1558	1559	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \		1560	1561	}	
	HOGCF48R	12	HOGCH23R	HOGCI28R	-	•								TOCOTION	VOI COOU		HOGCI42R	HOGOISSE	NCC COLUMN		HOGOP07R	HOGCEO, INCGODSERA 1561		
	HOGCE48	1	HOGCH23	HOGCI28											HOGCITO		1000140	H000011	HOGOS		TIOC DOLL	Section 1	HOGCFoo	

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Ala-71 to Trp-77, Pro-134 to Gly-146.	His-14 to Phe-20, Ser-37 to Gly-60.	Gly-20 to Leu-28,	Val-33 to Gly-41,	hr-43 to Gly-55,	al-58 to Arg-66.	Lys-7 to Thr-13,	Asp-24 to Thr-30,	ly-39 to Glu-52,	Leu-70 to Arg-76,	ne-87 to Tyr-92.	Lys-2 to Thr-8,	Asp-19 to Thr-25,	Gly-34 to Glu-47,	Leu-65 to Arg-71.	Asn-5 to Gln-12,	Gln-19 to Glu-25,	Arg-78 to Glu-99,	Thr-106 to Ala-111.	Arg-1 to Asn-8,	Phe-18 to His-24,	Asp-57 to Lys-62,	Lys-76 to Asp-81.	-	,	Thr-9 to His-19,	Leu-35 to Ala-46,
AY	3737 H S	3738 G	<u>> i</u>	<u> </u>	>	3739 L	<u>▼</u>	<u>o</u>	<u>.</u>	P	3740 L	<u> </u>	<u>છ</u>	L	3741 A	<u>5</u>	<u> </u>	Ŧ	3742 A	<u> </u>	<u> </u>	L	3743	3744	3745 TI	7
·	1 - 183	3 - 443				22 - 513					2 - 235				3 - 338			•	3 - 260				3 - 152	95 - 184	2 - 397	
	1562	1563				1564					1565				1566				1567				1568	1569	1570	
	HOGCT44R	HOGCV85R	_			HOGCV93 HOGCV93R					HOGCY12 HOGCY12R				HOGCY58R				HOGCY74R				HOGDDD29R	HOGDE77 HOGDE77R	HOGDG03R	
-	HOGCT44	HOGCV85				HOGCV93			- 1		HOGCY12				HOGCY58				HOGCY74				HOGDDD59	HOGDE77	HOGDG03	

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	0 0000	L0665: 3, H0682: 2, H0435: 2, L0751: 2,	H0663: 1, L0021: 1,	H064/: 1, L0800: 1, L0664: 1, H0683: 1, H0670: 1 and H0660: 1.																							
1. 65 to Dio.60	Gly-69 to Arg-74.		Phe-94 to Ser-99,	Thr-102 to Gly-107, Pro-113 to Ser-120.	60 to 61n-64	Cys-73 to Arg-88.		Pro-38 to Met-44.	Pro-6 to Tvr-17,	Val-39 to Gln-45.	Glu-3 to Glu-14.			Arg-9 to Val-1/.	Arg-9 to Arg-18,	Arg-27 to Arg-35,	Arg-47 to Asp-55,	Asp-78 to Lys-83.	_		Asp-89 to Thr-98.	_	Tyr-94 to Phe-99.		I en-3 to Arg-19.	+	
	40	3746			_	74/0	3748		T		3751	0,00	3/27	3753	3754				3755			3756	3757	十	十	+	3/60
	:	3 - 497	•			1 - 342	124 - 357	2 322	770	14-7-4/	2 200	5 - 477	3 - 335	22 - 492	48 - 470	!			1-417	ı		2:241	3 - 326	300	4	4	2 - 226
	,	1571	 -			1572	1573	1577		1575	1000	0/01	1577	1578	1				1580			1581	4		- 1		1585
		HOGDG76R	-			HOGDI44R	-1-	1.	J	~	٦,	HOGDPIOK	8	HOGDO95R	HOGDOS TOGODOS	VIOLUTION I			TOCHACO3P	מסטטון		TYOUR AND THE	HOGEN TO CH	HOGELUSOR	HOGEK25R	HOGEN30R	HOGEN55R
		HOGDG76				HOGDI44		HOGDI49	HOGD025 HOGD025F	HOGDO58		HOGDP10	110GD054	10000011	HOGDON	HOGDK/0				HOGDV93			HOGEAZ/	HOGED85	HOGEK25	HOGEN30	HOGEN55

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L0521: 1, L0662: 1 and H0435: 1.		H0295: 8, H0617: 7,	L0665: 7, H0435: 7,	H0592: 6, H0494: 6,	L0439: 6, L0751: 6,	S0360: 5, S0046: 4,	H0587: 4, H0012: 4,	H0264: 4, L0662: 4,	S0406: 4, H0586: 3,	H0620: 3, H0031: 3,	H0551: 3, S0440: 3,	S0002: 3, L0763: 3,	L0648: 3, L0649: 3,	L0774: 3, L0659: 3,	H0520: 3, H0547: 3,	H0689: 3, H0660: 3,	S0328: 3, S0330: 3,	L0748: 3, L0754: 3,	L0750: 3, H0294: 2,	H0650: 2, H0341: 2,	S0356: 2, S0132: 2,	H0370: 2, H0333: 2,	H0643: 2, L0623: 2,	H0706: 2, H0081: 2,	H0039: 2, H0553: 2,	H0181: 2, H0068: 2,	H0135: 2, H0040: 2,	H0646: 2, L0640: 2,
Gly-17 to Asn-27, Ser-51 to His-61.			•				<u>.</u>					•																
3761	3762	3763	:																								•	
3 - 212	2 - 223	2-376						-									•							-				
1586	1587	1588											•	-		- `						ż						
HOGEP46 HOGEP46R	HOGEP69R	HOGET60R	•																			-		-		<u>-</u>		
HOGEP46	HOGEP69	HOGET60	•														-											

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.0637: 2, L0641: 2,	0764: 2, L0804: 2,	.0657: 2, L0809: 2,	,0664; 2, H0519; 2,	10682: 2, FUO / U. 2, 10672: 2, L0744: 2,	L0740: 2, S0434: 2,	H0677: 2, H0483: 1,	0305: 1, H0638: 1,	0442: 1, S0358: 1,	0376: 1, S0410: 1,	H0411: 1, H0550: 1,	(0455: 1, H0600: 1,	H0101: 1, H0575: 1,	[0590: 1, H0618: 1,	10052: 1, H0234: 1,	H0597: 1, H0121: 1,	C0041: 1, H0123: 1,	H0246: 1, H0290: 1,	10292: 1, H0428: 1,	H0622: 1, T0023: 1,	H0030: 1, H0316: 1,	H0059: 1, H0100: 1,	S0306: 1, T0090: 1,	H0649: 1, S0210: 1,	S0426: 1, L0646: 1,	L0765: 1, L0771: 1,	L0626: 1, L0768: 1,	L0364: 1, L0766: 1,
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L0375: 1, L0651: 1, L0378: 1, L0653: 1, L0655: 1, L0558: 1, L0635: 1, L0518: 1, L0383: 1, L0792: 1, L0663: 1, S0006: 1, H0696: 1, S0044: 1, S3014: 1, L0743: 1, L0752: 1, L0755: 1, L0731: 1, L0759: 1, H0216: 1, H0543: 1, S0424: 1 and H0506: 1.	AR096: 1, AR104: 1, AR055: 0, AR061: 0, AR060: 0, AR089: 0 S0250: 1 and H0435: 1.							H0637: 2, H0413: 1 nd H0648: 1.			
	8					-6, r-56.	-10.				r-24.
	54 Pro-8 to Asn-34, Ala-39 to Arg-53	55	90	89	69	70 Ser-1 to Gln-6, Ser-51 to Thr-56.	71 Gly-4 to Lys-10		73	74	75 Pro-17 to Ser-24.
	3 - 368 3764	\vdash	256 - 333 3766 52 - 144 3767	╁	3 - 170 3769	128 - 313 3770	9 - 194 3771	9 - 152 3772	126 - 260 3773	60 - 383 3774	14 - 205 3775
	3EU84R 1589		HOGEW72R 1591	2	HOHDIISR 1594	OHB44R 1595	OHC27R 1596	HOOHJ69R 1597	DHK47R 1598	OHL15R 1599	OHL19R 1600
·	HOGEU84 HOGEU84R	HOGEW58 HOGEW58	HOGEW72 HOGEW72		1	ноонв44 ноонв44К	HOOHC27 HOOHC27	он 69гноон	HOOHK47 HOOHK47	HOOHL15 HOOHL15	ноонгія ноонгія

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.0666:	805: 4,	664: 4	662: 3	L0005: 2,	318:2	024: 2	770: 2	766: 2	665: 2	0648: 2	731:2	0677: 2	0583:	0586:	0486:	H0156: 1	0040	6028:	0615:	3036: 1	10040:	0472:	.0761:	0771:	0803:	1, 1.0775:	L0655:	,0659:	H0144:
L0748; 11, L0666: 5,	.0750: 5, L0805: 4,	.0663: 4, L0664: 4,	.0439: 4, L0662: 3,	3,10	H0587: 2, H0318: 2,	H0052: 2, H0024: 2,	H0553: 2, L0770: 2,	.0648: 2, L0766: 2,	.0497: 2, L0665: 2,	H0519: 2, H0648: 2,	J0740: 2, L.0731: 2,	.0593: 2, H0677: 3	H0170: 1, H0583:	H0638: 1, H0586:	H0331: 1, H0486:	3: 1, H	H0036: 1, L0040: 1	0: 1, S	H0328: 1, H0615: 1,	r0023: 1, S0036: 1	H0591: 1, H0040:	F0067: 1, S0472:	H0529: 1, L0761: 1	.0764: 1, L0771:	0649: 1, L0803:	4: 1, L	6: 1, L	.0657: 1, L0659:	.0790: 1, F
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		Lys-1 to Lys-13.	Pro-10 to Ser-37,	Gln-41 to Thr-57,	Arg-62 to Thr-68,	Pro-80 to Lys-85,	019-22 to 019-100.				Glu-1 to Glu-11.									<u>~ 4</u> ,			
·	3777	3778	3779			-	2700	2/00	3781	3782	3783	3784	3785										
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HOOIA57R 1611 251 - 361	L0803: 14, L0439: 9, L0752: 9, L0666: 7, L0659: 5, S0328: 5, L0756: 5, L0731: 5, L0794: 4, S0412: 4, S0360: 3, H0634: 3,
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2000	<u>8</u>	H05	<u>101</u>	90H	S03	0H	S03	H06	H05	10	HOH	HOI	H03	H	S04	H	S01	HO	HO	Ĥ)H	H	H	H	SO	H	윤	出	의
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	Ile-1 to Lys-7,	Gly-13 to Lys-19,	Pro-59 to Arg-76.	Val-50 to Arg-56.	Leu-41 to Ser-52.							Asp-48 to Thr-54,	Glu-57 to Ser-63.															
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L0749: 4, L0750: 4,	L0755: 4, H0663: 3,	S0418: 3, H0619: 3,	L0717: 3, H0013: 3,	L0163: 3, H0553: 3,	H0090: 3, L0763: 3,	L0770: 3, L0766: 3,	L0774: 3, L0664: 3,	L0665: 3, H0519: 3,	H0521: 3, L0742: 3,	L0591: 3, S0276: 3,	H0171: 2, H0341: 2,	S0356: 2, S0360: 2,	H0393: 2, S0300: 2,	H0156: 2, H0150: 2,	H0011: 2, H0687: 2,	S0003: 2, H0328: 2,	L0483: 2, H0634: 2,	H0494: 2, H0560: 2,	H0561: 2, H0646: 2,	L0638: 2, L0646: 2,	L0374: 2, L0764: 2,	L0776: 2, L0655: 2,	L0657: 2, H0520: 2,	H0660: 2, H0134: 2,	L0605: 2, L0593: 2,	H0542: 2, S6024: 1,	H0583: 1, S0116: 1,	H0483: 1, S0420: 1,	S0358: 1, S0408: 1,
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									Ala-2 to Asp-10.			Arg-11 to Pro-22, Asn-42 to Lys-48.				Pro-8 to Phe-18.
		3800	3801			3802	3803	3804	3805	3806	3807	3808				3809
		93 - 248	160 - 306			77 - 265	78 - 290	1 - 237	2 - 340	318 - 560	317 - 463	106 - 267				147 - 479
		1625	1626			1627	1628	1629	1630	1631	1632	1633				1634
		HOOJP58R				HOOJR72R	HOOJS84R	HOOJT32R	HOOJT65R	HOOJV78R	HOOJX42R	HOOJX80R				HOOJY17R
		HOOJP58	HOOJR60			HOOJR72	HOOJS84	HOOJT32	HOOJT65	HOOJV78	HOOJX42	HOOJX80				HOOJY17

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	Gln-1 to Cys-7, Asn-14 to Lys-Arg-29 to Leu-	Glu-46 to Leu-34.	Ala-1 to Pro-6, Ala-22 to Glu-27,	Glu-36 to Ser-45,	Glu-62 to Gly-67,	Gln-74 to Phe-83,	Thr-90 to Ser-96,	3													•				
	Gln-1 to Cys-7, Asn-14 to Lys-22, Arg-29 to Leu-34,	Glu-46	Ala-1 to Ala-22	Glu-36	Gln-62	Gln-74	Thr-90	7712-10																	
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		Asp-1 to Lys-22.			-			•				-	•		-		Pro-20 to Gly-38.		Gly-69 to Glu-78.		
-	3814	3815	3816														3817	3818	3819		
	107 - 367	118 - 453	210 - 344	-													14 - 196	107 - 571	207 - 461		
			1641														1642	1643	1644		
	HOOKK83 HOOKK83R	HOOKN43R	HOOKN71 HOOKN71R	_												•	HOPJA09R	HOPJF95R	HOPJG60R		
	HOOKK83	HOOKN43	HOOKN71														HOPJA09	HOPJF95	HOPJG60		

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777: 3, L0759: 3, 777: 2, L0809: 2, 7758: 2, H0341: 1, 9484: 1, S0358: 1, 9444: 1, S0360: 1, 5026: 1, S0278: 1, 0156: 1, H0102: 1, 0616: 1, H0102: 1, 0770: 1, L0766: 1, 0804: 1, L0657: 1, 0789: 1, H0684: 1, 10696: 1, S0044: 1, 0756: 1 and L0780: 1.	HO520: 18, S0344: 12, L0731: 11, T0006: 9, L0794: 9, L0753: 8, L0593: 8, S0278: 6, L0809: 6, H0135: 5, L0747: 5, H0065: 5, L0743: 5, L0747: 5, H0061: 3, H0594: 3, H0661: 3, H0666: 3, L0750: 3, L07
20777: 3, L0759: 3, L0774: 2, L0809: 2, L0758: 2, H0341: 1, S0444: 1, S0360: 1, S026: 1, S0278: 1, H0156: 1, H0102: 1, L0770: 1, L076: 1, L0789: 1, L0657: 1, L0789: 1, H0684: 1, L0789: 1, H0684: 1, L0789: 1, H0684: 1, L0756: 1 and L078: 1, L0756: 1, L0789: 1, H0684: 1, L0780:	H0547: 4 H0520: 18 L0731: 11, L0794: 9, L0593: 8, L0665: 5, L0747: 5, S0328: 4, H0661: 3, H0661: 3, H0658: 3 H0658: 3 H0658: 3 H0658: 3 H0658: 3 H0658: 3 H0658: 3
HHHW WALL	Ser-38.
	Pro-4 to Ser-29, Pro-31 to Ser-38.
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0: 2, H	2, S	9: 2, L(7:2, L	34: 2, H	19: 2, St	8: 2, L(.0595: 2, H0294: 1	H0656: 1, H0484:	59: 1, H	H0619: 1, H0586:	H0333: 1, H0427:	H0052: 1, H0251:	H0546: 1, H0545:	H0012: 1, H	H0181: 1, H0606:	H0334: 1, H0646:	S0144: 1, S0426:	.0762: 1, L	.0763: 1, L.0637:		.0803: 1, L0804:	Ξ,	0805: 1, L0629:	.0558: 1, L	.0783: 1, L0791:	H0689: 1, H0648:	S0330: 1, S0332: 1	36: 1, H	8: 1, L(
H005	H045	J_076	1051	H068	H053	L074	1059	H065	H066	H061	H033	H005	H054	H001	H018	H033	S014	1076	1076	L0662: 1	F080	L0774:	1080	1.055	1.078	H068	S033	H043	S002
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	Ala-83 to Arg-91, Trp-99 to Tyr-105, Tyr-123 to Glu-128.	Gly-2 to Ala-13.																							
	3821	3822	3823	3824																					
	3 - 542	166 - 597	321 - 464	148 - 285																					
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S004	H03,	H04,	H06	L052	1.077	L0517: 2,	467	104	S028	H066	S035	H032	H061	S030	S602	H05	H05	, H0590:	, H03	H05	H01	, S00	S038	, H0328:	H00	H00	H01	75	田田
H0036: 2, S0049: 2,	10052: 2, H0327: 2,	10687: 2, H0428: 2,	I0553: 2, H0644: 2,	10038: 2, L0520: 2,	.0762: 2,	.0653: 2,	30044: 2, L0442:	H0583: 1, L0443:	0: 1,	00:1,	S0348: 1, S0356: 1,	S0444: 1, H0329:	77: 1,	\$6026: 1,	.0717: 1, S6022: 1	50: 1,	HO587: 1, H0599:	H0042: 1,	H0318: 1, H0309:	45: 1,	H0564: 1, H0123: 1	H0019: 1, S0050:	.0163: 1, S0388: 1	10356: 1,	10688: 1, H0039:	70023: 1, H0031:	H0111: 1, H0169: 1	S0364: 1, L0455: 1	H0135: 1, H0163: 1
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	Ala-1 to Gly-11,	
	3825	3826
	6 - 497	101 - 250
	1650	1651
	HOPJH65R	HOPJM63R
	НОРЈН65	HOPJM63

H0039: 1, H0622: 1, S0464: 1, L0803: 1, H0684: 1 and H0670: 1.	L0665: 7, L0747: 7, L0665: 6, L0766: 6, L0655: 5, L0659: 5, S0126: 5, L0717: 4, H0370: 4, L0763: 4, L0770: 4, L0646: 4, H0521: 4, L0748: 4, S0360: 3, H0009: 3, S0142: 3, S0344: 3, L0771: 3, L0774: 3, L0771: 3, L0774: 3, L0775: 3, L0663: 3, L0751: 3, L0760: 3, H0265: 2, H0657: 2, S0418: 2, S0420: 2, S0418: 2, S0420: 2, S0418: 2, L0667: 2, L0372: 2, L0667: 2, L0372: 2, L0667: 2, L0372: 2, L0666: 2, L0748: 2, L0666: 2, L0748: 2, L0666: 2, L0748: 2, L0666: 2, L0748: 2, L0665: 2, L0748: 2, L0659: 2, L0748: 2, L0659: 2,
Lys-31 to Gly-36. H	Phe-15 to Ser-22. The ser-22 to Ser-22. The ser-22 to Ser-22 to Ser-22. The ser-22 to Ser-22 to Ser-22 to Ser-22.
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H0556: 1, H0295: 1,	S0114: 1, H0656: 1,	.0778: 1, S0282: 1,	H0255: 1, H0661: 1,	10402: 1, S0354: 1,	50358: 1, S0376: 1,	H0351: 1, S0222: 1,	HO586: 1, HO586: 1,	H0333: 1, H0250: 1,	T0082: 1, H0318: 1,	H0421: 1, H0052: 1,	.0738: 1, H0327: 1,	H0051: 1, H0083: 1,	50003: 1, S0022: 1,	S0214: 1, H0092: 1,	F0023: 1, L0483: 1,	r0006: 1, H0030: 1,	H0628: 1, H0181: 1,	H0169: 1, H0090: 1,	H0058: 1, L0351: 1,	F0041: 1, H0494: 1,	S0015: 1, L0065: 1,	S0426: 1, L0640: 1,	L0769: 1, L0772: 1,	L0641: 1, L0374: 1,	L0764: 1, L0773: 1,	L0648: 1, L0626: 1,	L0768: 1, L0649: 1,	L0784: 1, L0378: 1,	L0806: 1, L0776: 1,
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	o Ala-6
	Arg-5 to His-22, Ser-59 to Ala-69.
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	HOPJX48R
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L0663: 2, L0664: 2, L0665: 2, H0684: 2, L0741: 2, L0743: 2, H0170: 1, H0171: 1,	H0265: 1, H0583: 1, H0657: 1, H0656: 1, L0470: 1, H0254: 1, H0664: 1, H0662: 1,	S0420: 1, S0408: 1, H0393: 1, H0392: 1, H0613: 1, H0333: 1,	L0622: 1, H0101: 1, H0250: 1, H0427: 1,	50321: 1, H0255: 1, 50346: 1, H0421: 1,	H0620: 1, H0024: 1,	H0428: 1, H0553: 1, H0169: 1, H0674: 1,	S0366: 1, H0163: 1, H0040: 1, H0616: 1.	H0063: 1, H0026: 1,	H0517: 1, L0640: 1,	0761: 1, L0773: 1,	.0559: 1, L0659: 1,	.0790: 1, H0144: 1,	H0709: 1, H0696: 1,	H0187: 1, L0744: 1,
L0663 L0665 L0741 H0170	H0265 H0657 L0470 H0664	S0420 H0393 H0613	L0622 H025	1002 8034	H062	H042	S036	H006	H051	1,07	5507	201	COH LOH	HOI

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S0031: 1 and H0543: 1.	-		AR096: 4, AR089: 3,	AR061: 1, AR055: 1,	AR060: 1, AR104: 1	H0684: 2 and L0731:	1.							-	-				-			H0494: 6, H0556: 5,	H0265: 3, H0656: 3,	H0618: 3, H0295: 2,	H0255: 2, H0392: 2,	H0575: 2, H0253: 2,	H0052: 2, H0024: 2,	H0617: 2, H0087: 2,
		Pro-35 to Arg-48.	Arg-19 to Tyr-24,	Asp-75 to Glu-87,	Asn-102 to Cys-109, AR060: 1, AR104:	Ser-113 to Lys-118,	Tyr-154 to Ser-169.	Leu-39 to Ser-51.	Ser-50 to Pro-57.	Asn-80 to Gly-86.	Pro-6 to Thr-34,	Asp-57 to Gly-62,	Gly-77 to Glu-83,	Lys-108 to Asp-116,	Met-145 to Arg-159.	Ser-72 to Ala-78,	Asp-94 to Gly-100,	Thr-104 to Asp-111,	Ser-117 to Arg-133,	Glu-138 to Lys-144,	Lys-158 to Asp-165.	Ser-33 to Pro-42,	Lys-53 to Cys-61.					
	3829	3830	3831		•			3832	3833	3834	3835					3836						3837			,			
	273 - 395	2 - 163	2 - 676			,	,	26 - 232	3 - 191	1 - 456	1 - 579		,			1 - 663						3 - 455				•		
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	HOPJY50R	HOPJZ59R	HOPKA06R	•				HOPKC65R	HOPKD31R	HOPKG16R	HOPKG47 HOPKG47R					HOPKG83R			_			HOPKK38R						
	HOPJY50	HOPJZ59	HOPKA06					HOPKC65	HOPKD31	HOPKG16	HOPKG47					HOPKG83						HOPKK38						

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H0547: 2,	L0744: 2,	H0136: 2,	H0224: 1,	H0585: 1,	10341: 1,	H0483: 1,	S0418: 1,	30408: 1,	30045: 1,	H0550: 1,	H0587: 1,	, H0643: 1,	H0559: 1,	H0486: 1,	F0048: 1,	H0309: 1,	H0081: 1,	H0012: 1,	H0051: 1,	H0071: 1,	H0354: 1,	H0266: 1,	H0271: 1,	H0428: 1,	H0181: 1,	H0090: 1,	H0488: 1,	H0100: 1,	S0440: 1,
L0565: 2, H0547: 2,	H0521: 2, L0744: 2,	L0439: 2, H0136: 2	H0543: 2, H0224: 1	H0225: 1,	S0114: 1, H0341:	H0484: 1, H0483:	H0671: 1,	S0354: 1, S0408: 1,	S0007: 1, S	H0261: 1,	H0586: 1, H0587:	H0497: 1,	H0257: 1, H0559:	H0485: 1, H0486: 1	S0346: 1, T0048: 1	T0071: 1, H0309:	H0041: 1, H0081: 1	H0050: 1,	H0620: 1, H0051: 1	S0388: 1, H0071:	H0083: 1, H0354:	H0594: 1, H0266: 1	H0179: 1,	H0292: 1,	H0424: 1, H0181:	H0135: 1,	H0063: 1,	H0433: 1, H0100: 1,	T0041: 1, S0440:
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	HOPKN14R	HOPKN67R	HOPKO04 HOPKO04RA 1665														-			
	HOPKN14	HOPKN67	HOPKO04																	

H0412: 1, H0561: 1, S0150: 1, S0002: 1, H0529: 1, L0638: 1, L5565: 1, L0372: 1, L0649: 1, L0388: 1, L0789: 1, L0791: 1, H0519: 1, S0380: 1, H0522: 1, H0436: 1, L0747: 1, L0752: 1, L0757: 1, L0599: 1, H0542: 1, H0543: 1 and					L0439: 15, L0666: 8, S0222: 6, L0794: 5, L0638: 3, L0805: 3, L0664: 3, L0758: 3, L0770: 2, L0438: 2, H0624: 1, H0170: 1, S6024: 1, H0052: 1, H0672: 1, S6028: 1, H0572: 1, S6028: 1, H0572: 1, S6028: 1, H0572: 1, S6028: 1,
			Ser-19 to Arg-29, Asp-37 to Val-43, Ala-75 to Leu-80, Thr-96 to Arg-111.		
·	3841	3842	3843	3844	3845
·	1 - 405	282 - 536	19 - 402	1-519	3 - 173
	1666	1667		1669	
	HOPKO61R	HOPKP45R	HOPKQ20R	HODKO89R	HOPKRS6R
	HOPKO61	HOPKP45	НОРКQ20	1100770071	HOPKR56

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H0688: 1, H0616: 1, L0769: 1, L0768: 1, L0803: 1, L0774: 1, L0776: 1, L0659: 1, L0792: 1, H0519: 1, H0684: 1, L0745: 1, H0684: 1, L0745: 1, L0747: 1, L0756: 1, L0731: 1, L0759: 1 and L0731: 1, L0759: 1 and			L0794: 16, L0809: 6, S0360: 4, H0617: 3, L0803: 3, S0114: 2,	H0494: 2, L0375: 2,	L0731: 2, L0588: 2,	H0556: 1, S0134: 1, H0484: 1, S0420: 1	H0637: 1, H0441: 1,	H0370: 1, H0438: 1,	H0592: 1, T0109: 1,	H0009: 1, H0081: 1,	H0271: 1, H0428: 1,	S0368: 1, H0424: 1,	H0181: 1, H0124: 1,	S0438: 1, S0440: 1,
		Gly-1 to Val-7.	Pro-7 to 1 rp-14.											
	3846	3847	3848 848						•				•	
	2 - 274	2 - 136	237 - 551											
	1671	1672	1673											
	HOPKU33 HOPKU33R	HOVBH23R	HOVBK37R		:						•			
	HOPKU33	HOVBH23	HOVBK37											•

S0144: 1, S0142: 1, S0344: 1, L0637: 1, L5566: 1, L0761: 1, L0764: 1, L0548: 1, L0766: 1, L5574: 1, L0805: 1, L0783: 1, L0805: 1, L0783: 1, L0647: 1, L0789: 1, H0547: 1, H0670: 1, H0560: 1, H0648: 1, S0380: 1, H0648: 1, S0380: 1, H0521: 1, L0788: 1, L0779: 1, L0789: 1, L0779: 1, L0780: 1, L0757: 1,				S0360: 2, H0497: 2,	H0333: 1, H0004: 1, H0428: 1, H0674: 1, S0422: 1, L0763: 1,	L0803: 1, H0648: 1 and	30210: 1:			
SSZZZZZZZZ	His-7 to Lvs-20.	Glu-13 to Trp-32,	FTO-54 to visit				[C]	Gly-1 to Glu-7, Gly-29 to Ala-36.		Lys-63 to Ser-69,
	20.40	+	1300	1000	3852			3853	3854	3855
	700	144 - 404	-	41 - 349	342 - 524			2 - 127	395 - 580	3 - 578
	1	1675		1676	1677			1678	1679	
		HOVBK49R HOVBS15R		HOVBX41R	HOVCL23R			HOVCN03R	AACOOMOTE A	HOVCU89R
		HOVBK49 1		HOVRX41				HOVCN03		HOVCO26

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		·											
		H0615: 2, L0748: 2, H0656: 1, S0007: 1, H0486: 1, H0041: 1, H0355: 1, S0250: 1, S0003: 1, H0628: 1, H0428: 1, H0622: 1, T0042: 1, L0768: 1, L0657: 1, L0526: 1, H0702: 1, L055: 1, H0745: 1, L0749: 1, L0779: 1, L0588: 1,	LOCOGO, 1 8110 LOCOSC, 1.										
Tyr-95 to His-102,	Ser-15 to Tyr-21,			Ile-32 to Arg-37.						Asp-1 to Lys-6.			Pro-1 to Tyr-8,
	3856	3857	3858	3859	3860	3861	3862	3863	3864	3865	3866	3867	3868
	1 - 471	2 - 226	333 - 551	53 - 220	110 - 289	34 - 204	.1 - 270	37 - 498	24 - 257	253 - 393	92 - 184	74 - 154	3 - 365
	1681	1682	1683	1684	1685	1686	1687	1688	1689	1690	1691	1692	1693
-	HOVCV33 HOVCV33R	HOVCW45 HOVCW45R	HOVCY68 HOVCY68R	HOVCZ45R	HOVDB43R	HOVDB61R	HOVDB62R	HOVDB65R	HOVDD26R	HOVDE46R	HOVDG60R	HOVDG71R	HOVDH09R
	HOVCV33	HOVCW45	HOVCY68	HOVCZ45 HOVCZ45R	HOVDB43 HOVDB43R	HOVDB61	HOVDB62 HOVDB62R	HOVDB65 HOVDB65R	HOVDD26	HOVDE46 HOVDE46R	HOVDG60	HOVDG71 HOVDG71F	ноурно ноурноя

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		L0794: 10, L0803: 7, .0747: 6, H0657: 5,	H0597: 4, H0617: 4,	.0764: 4, L0809: 4,	0758: 4, L0759: 4,	0800: 3, L0804: 3,	,0775: 3, H0659: 3,	H0543: 3, H0231: 2,	.0471: 2, H0673: 2,	H0674: 2, L0763: 2,	.0770: 2, L0769: 2,	.0768; 2, L0649: 2,	.0375: 2, L0776: 2,	10658: 2, L0750: 2,	.0779; 2, L0755; 2,	.0731: 2, L0757: 2,	.0596; 2, L0588; 2,	S0026: 2, H0170: 1,	H0483: 1, H0661: 1,	H0402: 1, H0459: 1,	S0360: 1, H0639: 1,	L0717: 1, H0392: 1,	H0607: 1, H0333: 1,	H0486: 1, H0318: 1,
Gin-14 to Gin-19, Arg-25 to Giu-37, Asn-44 to Arg-50, Giu-67 to Leu-75, Thr-105 to Gly-114.		1)1	H	<u>a</u>	<u> </u>	7	<u>a</u>	<u>E</u>	<u></u>	<u>.=</u>	<u></u>		브.	<u> </u>		<u></u>		<u> </u>	_					
	3869	3870																						
	43 - 348	3 - 233																			•			
	1694	1695																						
·	HOVDH75R	HOVDU91 HOVDU91R															-							
	2700770175	HOVDU91											-								·			

H0251: 1, H0184: 1, H0045: 1, H0014: 1, S0388: 1, S0051: 1, H0594: 1, H0188: 1, H0328: 1, H0428: 1, H0181: 1, H0428: 1, H0181: 1, H0642: 1, H0413: 1, H0561: 1, S0372: 1, H0647: 1, S0144: 1, S0002: 1, H0026: 1, L074: 1, L0627: 1, L0648: 1, L0659: 1, L066: 1, L0659: 1, L0666: 1, L0663: 1, L0665: 1, L0663: 1, L0666: 1, L0663: 1, L0669: 1, L0751: 1, L0777: 1, L0751: 1, L0777: 1, L0752: 1, H0445: 1,	L0752: 3, L0731: 3, L0747: 2, L0753: 2, H0265: 1, S0116: 1, S0420: 1, H0428: 1, L0769: 1, L0767: 1,
	lle-12 to Gln-17.
	3871
	60 - 293
	1696
	HOVDV70 HOVDV70R
	HOVDV70

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L0766: 1, L0775: 1,	L0776: 1, H0555: 1, L0742: 1, L0748: 1, L0757: 1 and L0759: 1.	H0428: 1 and S3014: 1.	H0428: 2				S0474: 66. L0439: 16,	H0547: 15, L0748: 15,	L0731: 10, S0360: 9,	L0770: 9, S0408: 8,	S0358: 7, L0771: 7,	L0775: 7, L0747: 6,	L0592: 6, H0428: 5,	S0438: 5, L0659: 5,	L0664: 5, L0665: 5,	L0752: 5, L0757: 5,	H0506: 5, S0007: 4,	H0549: 4, S0440: 4,	L0769: 4, L0768: 4,	L0774: 4, L0663: 4,	H0519: 4, S0152: 4,	1 0750: 4, L0779: 4,	1.0755: 4, L0759: 4,	1,0591; 4, L0595; 4,	S0444: 3, H0510: 3,	H0615: 3, H0551: 3,	
		Tvr.22 to Ala-28.	Gly-40 to His-48.		Ala-20 to Gln-25.	Ser-7 to Trp-13,	Fne-19 to Ata-20.																		-		
		3077	3873	3874	3875	3876		3877						,													-
		1000	99-257	247 - 396	169 - 363	410 - 589		3 - 143																			
			1697					1702																			
			HOVDW60 HOVDW60R	HOVDZ66 HOVDZ66K	HOVEA43R	HOVEE84R		HOVEE92R																			
·		-	HOVDW60	HOVDZ66.	HOVEA43	HOVEB25	100 L	HOVEE92	No.																		

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H0100: 3, L0598: 3,	.0638: 3, LU/66: 3,	50380: 3, L0594: 3,	H0341: 2, S0420: 2,	S0442: 2, H0722: 2,	.0717: 2, H0550: 2,	S0222: 2, H0587: 2,	H0574: 2, H0632: 2,	H0013: 2, H0575: 2,	H0581: 2, H0544: 2,	H0012: 2, H0687: 2,	H0628: 2, S0036: 2,	H0509: 2, L0637: 2,	.0764: 2, L0773: 2,	.0803: 2, L0375: 2,	.0806: 2, L0783: 2,	.0666: 2, H0520: 2,	10684: 2, H0659: 2,	H0521: 2, L0745: 2,	.0756: 2, S0436: 2,	.0596: 2, L.0593: 2,	.0362: 2, S0194: 2,	H0543: 2, H0624: 1,	H0170: 1, H0171: 1,	H0717: 1, H0657: 1,	H0671: 1, H0663: 1,	H0664: 1, H0459: 1,	S0356: 1, S0410: 1,	H0637: 1, S0476: 1,	H0645: 1, S6026: 1,
OH.	3		OH.	20 <u>%</u>	2	30X	OH HO	OH OH	OH	OH	OH OH	<u>en</u>	<u>ra</u>	<u>3</u>	<u>3</u>	<u>3</u>	<u> </u>	OH HO	<u>3</u>	3	3	HO	OH.	OH .	HOH	OH)	SO	HO	HO
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9: 1, 6: 1	1: 1,): 1,	8: 1,	6: 1,	3: 1,	1: 1,	4: 1,	l: 1,	56: 1,	9: 1,	16: 1,	6: 1,	40: 1,	18: 1,	74: 1,	0: 1,	1,	55: 1,	73: 1,	46: 1,	52: 1,	51: 1,	76: 1,	57: 1,	84: 1,	45: 1,	92: 1,	38: 1,
S0300: 1, H0369: 1	H0497: 1, H0331: 1	L0021: 1, S0049: 1,	H0184: 1, L073	H0545: 1, H004	H0563: 1, H012	H0050: 1, L047	H0620: 1, H0014: 1,	S0388: 1, S005	H0356: 1, H026	S0003: 1, H003	H0622: 1, T0006: 1,	H0617: 1, S036	H0090: 1, H0040: 1	H0616: 1, S0038:	T0041: 1, H0494: 1,	S0014: 1, S015	L0369: 1, L06	L0763: 1, L550	1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1	L0372: 1, L06	L0521: 1, L06	L0804: 1, L0651: 1	L0652: 1, L07	1.0527: 1, 1.06	L0526: 1, L0384: 1	L0809: 1, L0545:	L0790: 1, L07	H0144: 1, L0438:
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H0593: 1, S0126: 1, H0682: 1, H0660: 1, H0648: 1, S0328: 1, H0539: 1, S0378: 1, S0350: 1, H0696: 1, C0740: 1, L0777: 1, L0758: 1, S0434: 1, H0217: 1, S0192: 1,		L0750: 11, L0740: 7, L0747: 5, H0341: 4,	L0757: 4, H0641: 3,	L0766: 3, L0748: 3,	L0754: 3, L0752: 3,	L0593: 3, H0171: 2,	S0134: 2, S0442: 2,	S0010: 2, H0428: 2,	H0538: 2, L0770: 2,	L0774: 2, L0776: 2,	L0517: 2, L0438: 2,	H0519: 2, H0684: 2,	L0777: 2, L0755: 2,	L0731: 2, L0758: 2,	S0026: 2, H0170: 1,	T0002: 1, S0114: 1,	S0116: 1, H0663: 1,	S0358: 1, S0360: 1,	S0007: 1, S0046: 1,
		Gin-4 to Gin-10, Lys-35 to Ala-60,	Pro-66 to Ser-79.														-		
	3878	3879																	
	288 - 497	1 - 393																	
·	1703	1704		_															
,	HOVEF34R	HOVEF81R									,								
	HOVEF34	HOVEF81																	

H0411: 1, H0592: 1, H0486: 1, H0013: 1, L0022: 1, H0575: 1, H0581: 1, H0052: 1, H0620: 1, H0012: 1, H0620: 1, H0034: 1, H00398: 1, S00346: 1, H00398: 1, S00346: 1, L0475: 1, L0520: 1, L06475: 1, L0520: 1, L0662: 1, L0522: 1, L0662: 1, L0521: 1, L0662: 1, L0552: 1, L0662: 1, L0659: 1, H0682: 1, H0689: 1, H0682: 1, H0689: 1, H0682: 1, H0689: 1, H0670: 1, S0330: 1, L0605: 1, H0436: 1, L0605: 1, H0542: 1 and H0423: 1.		L0809: 8, L0794: 5, L0439: 5, L0758: 5, H0617: 4, L0438: 4, L0754: 4, L0731: 4,
罗马马巴巴巴巴巴巴巴巴	Ser-9 to Gly-17, Ser-20 to Pro-26, Arg-32 to His-41.	
	3880	3881
	3 - 290	160 - 468
	1705	1706
	HOVEG73R	HOVEJISR
	HOVEG73	HOVEJ15

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H0295: 3, L0776: 3, H0619: 2, H0441: 2, H0253: 2, H0545: 2, H0188: 2, H0687: 2,	H0428: 2, H0424: 2, H0213: 2, H0031: 2, L0769: 2, L0803: 2, I 0792: 2, H0539: 2,	L0748: 2, L0745: 2, L0747: 2, L0757: 2, H0717: 1, H0549: 1, S0222: 1, H0333: 1	H0318: 1, S0049: 1, L0738: 1, H0046: 1, H0024: 1, H0090: 1,	H0649: 1, S0142: 1, L0763: 1, L0642: 1,	L0764: 1, L0771: 1, L0649: 1, L0651: 1, r 0630: 1-1 0805: 1	L0790: 1, H0144: 1,	H0723: 1, H0682: 1, H0696: 1, H0555: 1,	S0028: 1, L0742: 1, L0744: 1 and L0752: 1.				•
							·		Ser-13 to Val-21, Thr-45 to Ser-50.			
									3882	3883	3884	3885
									60 - 209	146 - 265	1 - 168	1 - 102
					<u> </u>				1707	1708	1709	1710
			,		,				HOVELSIR	HOVEN79R	HOVEN86 HOVEN86R	HOVEP45R
									HOVEL51	HOVEN79	HOVEN86	HOVEP45

	·									
	L0803: 7, L0748: 4, L0439: 4, L0804: 3, L0750: 3, L0756: 3, H0024: 2, L0438: 2, L0747: 2, L0779: 2, H0661: 1, S0354: 1, H0661: 1, S0354: 1, H0441: 1, H0331: 1, H0013: 1, S0214: 1, H0428: 1, H0040: 1, L0770: 1, L0776: 1, L0809: 1, L0665: 1, L0809: 1, L076: 1, L0770: 1, L0749: 1, L0777: 1, L0749: 1, L0777: 1, L0780: 1,									
	3887 Lys-5 to Ser-18.	Val-2 to Ser-7.				Ser-5 to Gly-10, Glu-18 to Glu-32,	Ser-89 to Arg-96,	Pro-141 to Ala-151,	Ser-162 to Gln-169.	Ala-18 to Ser-24, Gly-55 to Thr-60.
3886	3887	3888	3889	3890	3891	3892				3893
2-244	175 - 405	3 - 152	20 - 130	2 - 355	1 - 396	1 - 576				1 - 186
1711	1712	1713	1714	1715	1716	1717				1718
HOVEW80 HOVEW80R		HOVFD79 HOVFD79R	HOVJF86RA	HOVJH27 HOVJH27RA	HOVJI53R	HOVJJ09R				HOVJR56R
HOVEW80	HOVEY58	HOVFD79	HOVJF86	HOVJH27	HOVJI53	HOVJJ09				HOVJR56

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			L0157: 2, H0620: 2, L0666: 2, S0001: 1, L0717: 1, H0549: 1, S0222: 1, H0581: 1, H0194: 1, H0015: 1, H0399: 1, H0271: 1, H0688: 1, H0428: 1, H0672: 1, L0637: 1,					
Tyr-44 to Asp-49, Pro-132 to Leu-139, Cvs-168 to His-174.	Ala-8 to Gln-14, Arg-40 to Glu-45.	Pro-47 to Lys-53, Leu-93 to Lys-102, Tyr-105 to Gln-110,	Arg-113 to Lys-124. Ala-1 to His-13, Leu-73 to Leu-79.	Ala-20 to Val-38, Cys-43 to Thr-60, Arg-91 to Lys-97.	Gln-15 to Leu-20, Pro-37 to Thr-44.	Cys-23 to His-29, Ser-74 to Glu-91.	Gln-8 to Tyr-14, Pro-20 to Thr-26.	
3894	3895	3896	3897	3898	3899	3900	3901	3902
49 - 684	74 - 301	199 - 570	1 - 366	2 - 487	. 1 - 168	2 - 406	323 - 511	2 - 586
1719	1720	1721	1722	1723	1724	1725	1726	1727
HOVJU75 HOVJU75R	HOVJW17 HOVJW17R	HOVJY68R	HOVKB02R	HOVKE66 HOVKE66R	HOVKF20R	HOVKG18R	HPAMB11R	HPAMB60R 1727
HOVJU75	HOVJW17	ноулу68	HOVKB02	HOVKE66	HOVKF20	HOVKG18	HPAMB11	HPAMB60

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L0439: 11, L0747: H0624: 6, L0759: 6,	20779: 5, L0758: 5,	,0754: 4, H0171: 3,	.0756: 3,	.0752: 3, L0755: 3	.0591: 3, LU6U8: 3	H0170: 2, H0650: 2,	H0341: 2, H0663: 2,	S0360: 2, S0003: 2,	.0769: 2, L0796: 2,	,0766: 2, L0649: 2,	,0806: 2, H0696: 2,	S0028: 2, L0740: 2,	0750; 2, L0777; 2,	S0436: 2, L0361: 2,	S0298: 1, S0418: 1,	S0408:	H0733: 1, H0734:	S0300: 1, S0220:	H0497: 1, H0333:	H0575:	H0036: 1, H0590: 1	H0596: 1, H0050:	.0471: 1, H0015: 1	S0388: 1, S6028: 1	H0266: 1, H0328: 1	1, H0708: 1,	H0591: 1, H0040: 1
<u>L0439: 11, L0747: 8, 10624: 6, L0759: 6, </u>	79: 5, L	54: 4, F	0666: 3, 1	752: 3, L	191: 3, L	170: 2, 1	341: 2,]	360: 2, S	769: 2, 1	766: 2,]	806: 2,]	028: 2,]	750: 2,	436: 2,	298: 1,	444: 1,	5733: 1,	300: 1,	3497: 1,	0270: 1,	0036: 1,	0596: 1	0471: 1,	0388: 1,	0266: 1	H0622: 1	10591: 1
	107		3	5	ဌ	OH.) E	<u>SO:</u>	<u>2</u>	<u>2</u>	2	SO		80	S	80	Ĭ	SC	Ĭ	Ĭ	Ī	H	7	<u>X</u>	프	正	프
Ser-21																					•						
Leu-12 to Ser-21.																											
3903 I																											
2-196											_																
-	<u></u>								_																. 	_	
1728																											
/B62R																											-
HPA							_	_												_			<u></u>				·.
HPAMB62 HPAMB62R																											
HPA																							_				

					58: 14,
H0087: 1, H0412: 1, H0059: 1, S0112: 1, H0625: 1, S0440: 1, S0150: 1, S0422: 1, L0598: 1, H0529: 1, L0646: 1, L0770: 1, L0776: 1, L0774: 1, L0774: 1, L0775: 1, L0669: 1, L0669: 1, L0669: 1, L0669: 1, L0669: 1, H0659: 1, L0775: 1, H06445: 1, S0024: 1, L0749: 1, L0757: 1, H05443: 1, S0276: 1, H0543: 1, H0623: 1, S0412: 1 and H0423: 1, S0412: 1 and H023: 1, S0412: 1 and H023: 1, S0412: 1 and H023: 1, S0412: 1 and H0293: 1, S041					S0360: 15, S0358: 14,
					Arg-1 to Gly-6,
	3904	3905	3906	3907	3908
	2 - 439	134 - 334	8 - 631	1 - 669	18 - 464
	1729	1730	1731	1732	1733
	HPAMB93R	HPAMC04R	HPAMC19R	HPAMC27R	HPAMC90 HPAMC90R
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Γ		H0670: 1	S0422: 9	S0114: 7	S0408: 7	H0486: 7	S0444: 6	S0126: 6	S0378: 6	S0027: 6	H0040:	L0517: 5	S0404: 5	H0423: ;	H0171:	S0116: 4	H0545:	H0674:	1.0763:	1.0775:	L0518:	L0519:	H0445:	H0685:	S0420:	S0132:-	H0013:	H0597:	H0616:
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S0390: 2, S0032: 2,	.0742: 2, L0750: 2,	.0588: 2, L0603: 2,	S0026: 2, H0170: 1	H0395: 1, H0713:	S0402: 1, S6024: 1	H0295: 1, H0294:	S0218: 1, S0110:	H0402: 1, H0305: 1	S0418: 1, H0722:	H0728: 1, S0046: 1,	S0476: 1, S0300: 1	10462: 1, S0222:	H0385: 1, H0370:	H0404: 1, H0592:	H0497: 1, H0333: 1	L0586:	.0060: 1, H0098: 1	H0318: 1, L0563: 1	H0581: 1, H0421:	H0251: 1, H0309:	H0596: 1, H0231:	H0150: 1, H0123: 1	H0620: 1, L0163:	S0051: 1, H0399:	H0354: 1, H0266: 1	S0334: 1, S0312:	S0214: 1, H0428: 1	.0483: 1, T0006: 1,	H0553: 1, H0606: 1
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	Glu-10 to Phe-17, Phe-25 to Leu-32, Thr-38 to Tyr-44, Gln-81 to Thr-88.	Gly-60 to Phe-66.
	3909	3910
	1 - 504	2 - 277
	1734	1735
	HPAMD56 HPAMD56R	HPAME35R
	HPAMD56	HPAME35

H0597: 1, H0545: 1, F0003: 1, H0510: 1, S0314: 1, H0039: 1, L0194: 1, T0086: 1, H0674: 1, H0708: 1, H0579: 1, H0708: 1, H0598: 1, H0634: 1, H0509: 1, S0344: 1, L075: 1, L0764: 1, L0659: 1, L0764: 1, L0659: 1, L0764: 1, L0657: 1, L0806: 1, L0657: 1, L0806: 1, H0657: 1, L0806: 1, L0657: 1, L0806: 1, L0657: 1, L0788: 1, L0657: 1, L0669: 1, H0670: 1, H0689: 1, H0670: 1, H0660: 1, H0670: 1, H0689: 1, H0670: 1, H0660: 1, H0670: 1, H0660: 1, H0670: 1, H0660: 1, H0670: 1, H0660: 1,	H0663: 1, L0438: 1 and H0682: 1.	
a part of the second of the se	Lys-1 to Leu-12.	Glu-1 to Gly-8.
	3911	3912
	184 - 492	19-513
	1736	1737
	HPAME58R	HPAMF16R
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L0752: 12, L0748: 8, L0766: 5, L0599: 5,	H0617: 4, L0764: 4, H0081: 3, L0774: 3,	L0749: 3, L0750: 3,	S0444: 2, S0408: 2, H0392: 2, H0150: 2.	H0494: 2, H0521: 2,	L0756: 2, L0758: 2,	H0265: 1, H0483: 1,	S0358: 1, L0021: 1,	H0108: 1, H0024: 1,	H0083: 1, S0366: 1,	H0090: 1, H0038: 1,	ئـ آ	L3905: 1, L0772: 1,	L0644: 1, L0645: 1,	3		L0659: 1, L0666: 1,	L0665: 1, L0438: 1,	H0519: 1, H0682: 1,	H0670: 1, S0188: 1,	L0740: 1, L0747: 1,	L0777: 1, L0759: 1,	H0445: 1, L0596: 1,	L0604: 1 and H0542: 1.
Ser-35 to Gly-41, Ser-71 to Met-80.		بس													•		-						
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		Lys-13 to Tyr-30,	Arg-52 to Lys-57,	Ser-72 to Thr-79,.	Thr-110 to Trp-115,	Leu-124 to Ala-130,	Leu-132 to Glu-138,	Asn-141 to Lys-162.					-					<u> </u>				<u></u>	. 	<u>Uj</u>	<u> </u>	1	Pro-18 to Lys-24.	Thr-27 to Lys-37.
3914	3915	3916							3917	3918	•																3919	3920
385 - 666	3 - 410	2 - 670							2 - 472	100 - 246							,	 									2 - 190	327 - 509
1739	1740	1741							1742	1743																	1744	1745
HPAMG44 HPAMG44R	HPAMG54R	HPAMIIIR							HPAMJ71R	HPAMQ47 HPAMQ47R			_														HPAMQ76 HPAMQ76R	HPAMT47R
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H065	H065	3, 1.075	3, 1.073), H025	2, H058	2, H05	2, L079	2, L07	2, S041	1, H06	1, S044	1, H03	1, H06	1, H03	1, H00	1, L01	1, H01	1, H05	1, 1,01	1, S0051:	30336: 1, S0214:	10428: 1, H0032: 1	H0166: 1, H0673: 1,	H0561: 1, S0440:	1, 1.07	J0803: 1, L0804:	1, TOC	: 1, HO	H0593: 1, H0690: 1
0744.4 H0656: 3.	0766.3 H0659:3.	10436: 3, L0754: 3,	L0747: 3, L0731: 3,	0026: 3	S0007: 2, H0587: 2,	.0586: 2, H0545: 2,	H0150: 2, L0794: 2,	H0144: 2, L0750: 2,	,0608: 2, S0412: 2,	H0170: 1, H0638: 1,	S0420: 1, S0442:	S0444: 1, H0329:	30045: 1, H0645: 1	H0586: 1, H0333:	H0574: 1, H0013:	10250:	H0581: 1, H0196: 1	H0263: 1, H0546: 1	H0457: 1, L0157: 1	.0471: 1	30336:	H0428:	H0166:	H0561:	L0763: 1,	L0803:	L0776: 1,	H0547	H0593
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	Lys-59 to Tyr-70, Pro-109 to Lys-115.	Gln-34 to Asn-39.	Gly-1 to Arg-13,	Glu-37 to Lys-42,	Gly-45 to Gln-64.																
	3921	3922	3923																		
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	Pro-1 to Cys-6, Ala-13 to Ala-19, His-22 to Lys-28, Ser-44 to Cys-53, Pro-56 to Gly-67.	Trp-19 to Ser-27, Pro-185 to Ser-195.	
	3924	3925	3926
	291 - 569	2 - 628	3 - 149
	1749	1750	1751
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H0388: 1, H0400: 1, S0036: 1, H0090: 1, H0551: 1, H0264: 1, H0412: 1, S0386: 1, T0042: 1, H0625: 1, L0625: 1, L0506: 1, L0517: 1, L0665: 1, L0517: 1, L0665: 1, L0517: 1, L0666: 1, L0664: 1, S0374: 1, H0647: 1, H0648: 1, H0658: 1, H0648: 1, H0658: 1, H0658: 1, H0658: 1, H0648: 1, S0146: 1, S0390: 1, L0747: 1, L0753: 1, S0242: 1, S0194: 1, S0242: 1, H0542: 1 and	S0424: 1.
	Ser-1 to Lys-25, Glu-71 to Ser-77, Glu-93 to Thr-99,
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							H0497: 5, S0002: 5,	H0529: 5, H0556: 4,	H0046: 3, L0455: 3,	T0042: 3, H0520: 3,	H0543: 3, H0265: 2,	H0657: 2, H0656: 2,	H0458: 2, S0358: 2,	H0486: 2, H0050: 2,	H0124: 2, H0591: 2,	H0494: 2, L0662: 2,	H0522: 2, L0740: 2,	L0751: 2, L0747: 2,	L0588: 2, L0605: 2,	H0667: 2, H0686: 1,	H0650: 1, H0664: 1,	H0125: 1, S0418: 1,	H0580: 1, S6014: 1,	H0257: 1, H0559: 1,	H0635: 1, H0427: 1,	H0575: 1, H0052: 1,	N0006: 1, L0471: 1,	H0373: 1, H0083: 1,
Arg-123 to His-134,	Pro-141 to His-146, His-198 to Ala-207.	Thr-23 to Gln-35,	His-37 to Val-51,	Arg-70 to Arg-81,	Gln-108 to Asp-114,	Met-130 to Ser-140.		7,		Val-105 to Leu-110,	Thr-139 to Thr-153, H0543: 3, H0265: 2,	Thr-165 to Lys-187,	Ala-198 to Leu-203, H0458: 2, S0358: 2,	Ser-211 to Lys-226.									-	•				
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		HPANAOSR					HPANA07R	W/OWNE III																				
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H0266: 1, S0250:	S0022: 1, H0428: 1	H0031: 1, H0383:	H0212: 1, S0366:	S0036: 1, H0040:	H0634: 1, H0561:	S0210: 1, L0770:		L0641: 1, L0644:	L0764: 1, L0776:	L0655: 1, L0807:	L0657: 1, L0665:	 · 🚅	ì 🛁	·	L0596: 1, L0485:	L0601: 1, S0276: 1	·	S0446: 1.		•							٠
	-				-														Gln-22 to Thr-31,	Leu-91 to Leu-104.	Gln-13 to Gly-28.		Leu-1 to Ser-8.		Lys-64 to Tyr-75,	Pro-114 to Lys-120,	Leu-160 to Lys-169,
																			3930		3931	3932	3933	3934	3935		
											٠								2-376		494 - 730	2 - 247	255 - 446	2-412	1-666		
															•				1755		1756	1757	1758	1759	1760		
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														•			•		HPANA28		HPANA87	HPANB32	HPANC01	HPANE49	HPANE52		

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	L0752: 10, L0593: 9,	L0775: 8, L0588: 8,	L0666: 7, L0665: 7,	L0592: 7, L0595: 7,	L0363: 6, L0774: 6,	L0747: 6, L0362: 6,	H0422: 6, L0455: 5,	L0659: 5, H0539: 5,	H0170: 4, S0360: 4,	L0657: 4, H0547: 4,	L0749: 4, L0779: 4,	L0757: 4, H0657: 3,	H0412: 3, L0520: 3,	L0761: 3, L0646: 3,	L0375: 3, L0651: 3,	L0783: 3, H0144: 3,	H0520: 3, H0659: 3,	H0670: 3, H0648: 3,	L0740: 3, L0750: 3,	L0756: 3, L0731: 3,	L0759: 3, L0591: 3,	L0608: 3, H0662: 2,	S0376: 2, H0637: 2,	H0619: 2, L0717: 2,	S0280: 2, H0318: 2,	T0010: 2, H0040: 2,	H0413: 2, L0351: 2,	H0646: 2, L0763: 2,
Tyr-172 to Gln-177, Pro-182 to Val-189.								•																				
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	1761			•							•																	
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.0521: 2, L0784: 2,	0655: 2, L0382: 2,	0809: 2, L0664: 2,	10682: 2, H0658: 2,	10672: 2, L0439: 2,	0755: 2, L0589: 2,	0604: 2, S0424: 2,	H0624: 1, H0686: 1,	H0685: 1, S0040: 1,	H0716: 1, H0341: 1,	H0580: 1, S0468: 1,	S6026: 1, H0351: 1,	H0392: 1, L0623: 1,	060: 1, H0069: 1,	H0427: 1, T0048: 1,	70071: 1, S0049: 1,	H0052: 1, L0471: 1,		S0003: 1, T0006: 1,	.0055: 1, H0169: 1,	S0036: 1, T0067: 1,	H0059: 1, H0494: 1,	H0560: 1, S0372: 1,	S0002: 1, L0371: 1,	.0637: 1, L0627: 1,	.0772: 1, L0764: 1,	.0662: 1, L0767: 1,	.0768: 1, L0649: 1,	4500: 1, L0650: 1,	_0378: 1, L0806: 1,
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	1 1-1															Ser-5 to Ser-14, Gln-21 to Gln-28.		-			· · ·								Pro-1 to Val-6,	
			_													3937	3038	0000									3939	3940	3941	
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	·															HPANJ67R		HPANKOIK									HPCOR29R	┥-		4
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		H0591: 62, S0242: 15,	L0754: 11, S0003: 9,	S0196: 8, L0766: 7,	H0144: 6, L0740: 6,	L0664: 5, S0420: 4,	H0393: 4, L0803: 4,	H0428: 3, H0509: 3,	H0641: 3, L0777: 3,	L0595: 3, L0362: 3,	S0360: 2, H0013: 2,	L0471: 2, L0163: 2,	H0355: 2, H0553: 2,	S0036: 2, L0662: 2,	L0804: 2, S0374: 2,	L0438: 2, S0126: 2,	L0748: 2, L0755: 2,	L0485: 2, S0412: 2,	H0341: 1, S0376: 1,	H0580: 1, S0045: 1,	H0331: 1, T0039: 1,	T0060: 1, H0427: 1,	L0021: 1, H0590: 1,	H0318: 1, H0194: 1,	H0563: 1, H0572: 1,	H0051: 1, H0083: 1,	S6028: 1, S0312: 1,
Asp-51 to Phe-56, Asp-83 to Gly-88, Ser-94 to Glu-102.		Ile-12 to Cys-18.								•										`							
	3942	3943															:										
	197 - 346	426 - 557										•							-							-	
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	HPCOK72R	HPCOK73 HPCOK73R																									
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S0214: 1, H0031: 1, H0628: 1, H0598: 1, H0616: 1, H0534: 1, H06412: 1, L0370: 1, H0494: 1, H0625: 1, S0440: 1, S0344: 1, L0646: 1, L0543: 1, L0646: 1, L0545: 1, L0775: 1, L0521: 1, L0775: 1, L0521: 1, L0775: 1, L0521: 1, H0519: 1, H0711: 1, H0519: 1, H0711: 1, H0519: 1, H0711: 1, H0519: 1, H0660: 1, L0759: 1, L0592: 1, L0594: 1, H0667: 1, L0594: 1, H0667: 1, H0423: 1, H0422: 1,						
	07 1 12 147	Lys-42 to Val-49.	Ala-3 to Gly-9.	Pro-28 to Asn-33,	Alg-45 to Ash-55.	
v.'	3944	3945	3946	it C	3948	
	1 - 108	33 - 179	3-335	24- 240	2 - 427	
	1769	1770	1771	1//2	1773	7,74
	HPCOK93R	HPCOL81R	HPCO090R	HPCOO95K	HPCOD3R	TI COI 4011
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	L0803: 8, L0777: 8,	L0794: 7, L0766: 7,	H0144: 5, L0438: 5,	H0547: 5, L0439: 5,	H0013: 4, L0779: 4,	H0441: 3, S0440: 3,	L0762: 3, L0771: 3,	L0662: 3, L0666: 3,	L0665: 3, L0748: 3,	L0752: 3, H0656: 2,	H0341: 2, S0418: 2,	S0356: 2, H0575: 2,	H0328: 2, H0615: 2,	H0040: 2, H0477: 2,	T0041: 2, L0598: 2,	L0763: 2, L0659: 2,	L0663: 2, L0664: 2,	H0659: 2, L0602: 2,	S0406: 2, L0755: 2,	L0759: 2, L0592: 2,	H0624: 1, H0170: 1,	H0171: 1, S0134: 1,	H0583: 1, H0650: 1,	H0657: 1, L0415: 1,	S0116: 1, H0661: 1,	H0664: 1, S0420: 1,	L0005: 1, S0444: 1,	S0360: 1, S0408: 1,	S0046: 1, S0476: 1,
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3949	3950			•		•												•							,				
417 - 548	2 - 259			_			•																-						***
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S0300: 1, L0717: 1,	H0411: 1, H0486: 1,	F0039: 1, H0318: 1,	H0581: 1, H0123: 1,	H0014: 1, H0354: 1,	1, S0003: 1,	H0628: 1, H0032: 1,	l, L0060: 1,	H0264: 1, H0412: 1,	i, H0625: 1,	H0131: 1, H0646: 1,	S0002: 1, L0770: 1,	l, L0761: 1,	l, L0764: 1,	L0649: 1, L0804: 1,	l, L0519: 1,	.0352: 1, H0519: 1,	S0126: 1, H0658: 1,	1, H0672: 1,	80328: 1, 80330: 1,	1, H0696: 1,	H0704: 1, S0027: 1,	1, L0754: 1,	1, L0758: 1,	1, L0591: 1,	S0011: 1, H0543: 1,	H0422: 1, H0506: 1 and	1.		
S0300: 1	H0411: 1	T0039: 1	H0581: 1	H0014: 1	H0179: 1	H0628:	S0036: 1	H0264:	L0564: 1	H0131:	80002: 1	1.0637:	T0800:	10649:	L0650:	10352:	S0126: 1	H0648:	50328:	H0518:	H0704:	L0743:	L0747:	80434:	S0011: 1	H0422:	H0721:		Ala-44 to Gly-50.
																												3951	3952
				-																								483 - 641	133 - 396
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	S0414: 16, S0360: 6,	S0358: 5, L0766: 5,	H0574: 4, H0521: 4,	L0754: 4, L0756: 4,	S0422: 3, L0770: 3,	H0547: 3, H0672: 3,	L0596: 3, H0251: 2,	H0615: 2, H0090: 2,	L0646: 2, L0662: 2,	L0776: 2, L0659: 2,	L0752: 2, L0758: 2,	\$0436: 2, L0595: 2,	H0624: 1, L0002: 1,	H0583: 1, H0650: 1,	S0116: 1, S0298: 1,	S0282: 1, S0418: 1,	H0722: 1, H0734: 1,	S0045: 1, S0300: 1,	L0717: 1, S0222: 1,	H0013: 1, L0021: 1,	H0706: 1, L0105: 1,	H0263: 1, L0041: 1,	H0057: 1, S0050: 1,	L0163: 1, H0051: 1,	S0388: 1, S0051: 1,	H0271: 1, S0003: 1,	H0644: 1, H0032: 1,	H0673: 1, H0379: 1,	H0272: 1, H0059: 1,
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H0130: 1, H0647: 1, S0210: 1, S0426: 1, H0529: 1, L0637: 1, L0372: 1, L0649: 1, L0775: 1, L0651: 1, L0806: 1, L0657: 1, L0518: 1, L0788: 1, L0665: 1, H0144: 1, L0665: 1, H0519: 1, H0689: 1, H0519: 1, H0689: 1, H0478: 1, H0659: 1, S0406: 1, H0659: 1, S0406: 1, H0727: 1, L0439: 1, L0731: 1, H0595: 1, S0434: 1, L0608: 1, H0667: 1, S0196: 1, H0622: 1, S0424: 1 and H0622: 1, S0424: 1 and				L0439: 3, L0438: 1, H0659: 1, H0648: 1 and S0330: 1.		H0551: 11, L0766: 10, L0794: 8, S0026: 7,
H0130: 1, 10372: 1, 10372: 1, 10372: 1, 10372: 1, 10372: 1, 10373: 1, 10373: 1, 10438: 1, 10436:	Lys-4 to Lys-16.		Asn-9 to Trp-14, Pro-20 to Arg-46.			L07
·	Н	3956	3957	3958	3959	7 3960
	151 - 657	3 - 356	158 - 418	294 - 473	3-314	308 - 427
	1780	1781	1782	1783	1784	1785
	HPCOV35R	HPCOV41R	HPCOX43R	HPCPB38R	HPCPD26R	HPCPE33R
	HPCOV35	HPCOV41	HPCOX43	HPCPB38	30 dayan	HPCPE33

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70: 6,	521: 6,	61: 4,	197: 4,	170: 3,	60:3	561: 3,	57: 3,	18: 2,	56: 2,	78: 2,	10: 2,	14: 2,	517: 2,	90: 2,	66: 2,	02: 2,	69: 2,	L0646: 2,	L0655: 2,	89: 2,	1,0664: 2,	L0758: 2,	H0668: 2,	123: 2,	42: 1,	57: 1,	02: 1,	76: 1,	139: 1,
6, 1.07	6, HO	5, H06	4, H0	4, H0	3, S03	3, H0	3, L07	2, S04	2, S03	2, S02	2, T01	2, S02	2, H06	2, H00	2, H03	2, S00	2, 1.03	2, L06	2, L06	2, L0789:		2, L07	2, H06	2, H02	1, S03	1, H06	1, H04	1, S03	1, H03
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S0194: 1, H0542: 1, H0422: 1 and H0506: 1.	H0659: 2, L0752: 2, L0595: 2 and H0686: 1					L0766: 10, L0754: 9,	S0358: 6, H0616: 6,	H0551: 5, L0439: 5.	H0657: 4, H0497: 4,	S0003: 4, H0038: 4,	S0422: 4, S0374: 4,	H0659: 4, S0045: 3,	H0644: 3, H0412: 3,	H0529: 3, L0662: 3,	L0805: 3, L0657: 3,	L0663: 3, H0521: 3,	H0624: 2, H0171: 2,	S0040: 2, H0638: 2,	S0418: 2, L0005: 2,	S0376: 2, H0546: 2,	H0046: 2, H0553: 2,	H0169: 2, H0591: 2,	H0413: 2, H0623: 2,	L0646: 2, L0771: 2,	L0803: 2, L0526: 2,	L0666: 2, L0664: 2,
		Pro-10 to Val-15, Ser-25 to Asn-32.			Pro-29 to Gln-38.	Gln-47 to Lys-55.																				
	3961	3962	3963	3964	3965	3966																				
	93 - 209	203 - 412	3 - 101	191 - 322	3 - 116	201 - 407									-		•									
	1786	1787	1788	1789	1790	1791		·																		
	HPCPH50R	HPCPH51R	HPCPH52R	HPCPH68R	HPCPU27R	HPCPW09 HPCPW09R						-											•			
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380: 2,	3777: 2,)242: 2,	0656: 1,	0341: 1,	0671: 1,	1, 1,	1132: 1,	0717: 1,	0574: 1,	0486: 1,	0013: 1,)474: 1,	0545: 1,	0373: 1,	0688: 1,	0483: 1,	0055: 1,	0674: 1,	0090: 1,	T0041: 1,	0561: 1,	0509: 1,	0646: 1,	3598: 1,	0369: 1,	0773: 1,	0774: 1,	0655: 1,	0659: 1,
S0330; 2, S0380; 2,	L0748: 2, L0777: 2,	L0758: 2, S(H0170: 1, H0656: 1	S0116: 1, H0341: 1	S0212: 1, H0671: 1	S0356: 1, SC	S0360: 1, S0132:	H0619: 1, L	S0278: 1, H	H0632: 1, H0486: 1,	T0114: 1, H	L0105: 1, S(T0110: 1, H	L0471: 1, H	H0290: 1, H	H0622: 1, L	H0628: 1, L	H0673: 1, H0674: 1,	H0068: 1, H	H0056: 1, T	H0560: 1, H0561:	S0440: 1, H0509:	S0150: 1, H0646:	S0144: 1, L	H0517: 1, L0369:	L0372: 1, L0773:	L0804: 1, L0774:	L0653: 1, L0655:	L0527: 1, L0659: 1
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L0518: 1, L0783: 1, L0809: 1, L0647: 1, H0144: 1, H0520: 1, H0648: 1, S0328: 1, S0152: 1, H0555: 1, S3014: 1, L0747: 1, L0749: 1, S0031: 1, L0688: 1, S0436: 1, L0608: 1, L0595: 1, S0011: 1, S0026: 1, H0136: 1, H0543: 1,	H0659: 2, L0438: 1 and L0439: 1.		H0521: 13, L0659: 10,	L0779: 10, H0551: 9,	S0360: 6, S0214: 6,	L0770: 6, L0649: 6,	L0666: 6, L0731: 6,	H0170: 5, H0638: 5,	S0376: 5, H0580: 5,	H0575: 5, H0553: 5,	S0210: 5, L0662: 5,	L0776: 5, L0657: 5,	H0547: 5, L0439: 5,	H0656: 4, S0358: 4,	H0676: 4, H0574: 4,	H0486: 4, H0318: 4,	S0003: 4, L0655: 4,
	His-25 to Lys-30.	Thr-19 to Leu-24.		•			-								,		
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.0663: 4, L0665: 4,	H0659: 4, L0756: 4	.0757: 4, H0445: 4,	H0542: 4, S0212: 3,	H0009: 3, LO157; 3,	.0471: 3, H0024: 3,	H0494: 3, L0763; 3,	.0637: 3, L.0646: 3,	.0764: 3, S0052: 3,	H0672: 3, L0748: 3,	J0750: 3, L0752: 3,	.0755: 3, L0608: 3,	S0026: 3, H0543: 3,	H0171: 2, H0556: 2,	H0657: 2, L0785: 2,	S0420: 2, L0005: 2,	H0156: 2, S0474: 2,	H0581: 2, H0251: 2,	H0545: 2, H0373: 2,	H0477: 2, H0412: 2,	H0413: 2, T0041: 2,	S0466: 2, H0641: 2,	S0144: 2, L0372: 2,	.0774: 2, L	.0783: 2, L(S0374: 2, L0438: 2,	H0658: 2, H0670: 2,	H0660: 2, H0539: 2	H0522: 2, H0436: 2,	J0742: 2, L0777:
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	3970	3971	$\neg \neg$	3973 Pro-3 to Cys-9, Asn-28 to Phe-37	3974	
	113 - 271	╁	7 1 - 345	3 2-229	9 259 - 408	
	HPCOOKER 1795	HPCOT27R	HPCOT88R 1797	HPCQW64R 1798	HPCQW77 HPCQW77R 1799	
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S0360: 2, L0769: 2,	.0649; 2, H0648; 2,		.0747: 2, L0777: 2,	J0755: 2, L0757: 2,		70049: 1, H0657: 1,	H0255: 1, H0638: 1,	S0408: 1, H0619: 1,	H0037: 1, H0052: 1,	.0471: 1, H0014: 1,	H0316: 1, S0438: 1,	.0520: 1, L0763: 1,	.0770: 1, L0637: 1,	.0761: 1, L0373: 1,			•	L0527: 1, L0542: 1,	.0382: 1, L0519: 1,	H0144: 1, H0682: 1,	H0659: 1, H0660: 1,	H0696: 1, H0555: 1,	S3012: 1, L0439: 1,	.0751: 1, L0754: 1,	.0745: 1, L0749: 1,	.0752: 1, L0759: 1,	H0542: 1, S0458: 1 and	S0384: 1.	
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	Ser-8 to Lys-21, Gln-47 to Glu-52, Are-72 to Are-87.	Met-4 to Lys-18, Gln-44 to Glu-49,	Asn-1 to Lys-20, Gln-46 to Glu-51, Arg-71 to Arg-86.	Asn-1 to Lys-21, Gln-47 to Glu-52, Arg-72 to Arg-87.															-	
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		Lys-37 to Arg-42, Ser-74 to Thr-81.	Arg-6 to Asp-15.			•													
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L0352: 1, H0670: 1, L0744: 1, L0740: 1, L0780: 1, L0759: 1 and H0506: 1.	L0803: 2, H0413: 1, L0650: 1, L0666: 1, H0689: 1, H0659: 1, H0658: 1, L0758: 1, L0759: 1 and S0242: 1.			-		L0439: 11, L0740: 10,	10748.7 10005.6	S0358: 6, L0471: 6,	S0214: 6, L0803: 6,	L0517: 6, L0666: 6,	L0759: 6, L0664: 5,	L0665: 5, S0126: 5,	H0574: 4, H0156: 4,	L0646: 4, L0771: 4,	L0766: 4, S0374: 4,	H0659: 4, L0754: 4,	L0749: 4, L0731: 4,	L0362: 4, H0170: 3,	H0341: 3, H0632: 3,	S0003: 3, H0413: 3,	H0529: 3, L0369: 3,
		Gly-4 to Lys-9.		Pro-7 to Ala-13.		Asn-27 to Lys-36.		-						-							-
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S04.	105	H06	H05	H03	S028	90H	800 <u>x</u>	H04	H01:	H02	S02	90H	<u>100</u>	90H	S03	<u>100</u>	HOS	90H	<u>1</u> 02	747	106	<u>2</u>	<u>10</u>	<u>1</u> 07	<u>8</u>	106	<u>106</u>	105	1.05
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	Asp-1 to Leu-6.		Gln-20 to Leu-25.	Asp-11 to Val-19, Cys-36 to Asp-58, Lys-102 to Lys-111, Ile-115 to Ala-125, Val-127 to Gln-132.
	3995		3997	
	57 - 209	1 - 99	3 - 287	2 - 412
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HOOLI	L0521	10659	S0374	H0423	H0716	S0376	S0474	H0553	H0412	L0651	L0438	H0435	S0027	H0665	H0543	T0049	S0442	S0410	S0476	H0427	H0706	H0581	H0530	T0010	H0615	H0617	H0673	H0038	S0150
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38: 2.	64: 2,	.0768: 2,	66: 2,	L0806: 2,	L0809: 2,	L0665: 2,	25: 2,	26: 2,	582: 2,	328: 2,	696: 2,	128: 2,	L0755: 2,	1.0759: 2,	436: 2,	L0599: 2,	136: 2,	624: 1,	686: 1,	430: 1,	650: 1,	282: 1,)661: 1,)662: 1,	, H0300: 1,	720: 1,	0580: 1,	046: 1,	278: 1,
h 0598· 2 1 0638: 2.	1.0646; 2, L0764; 2,	L0773: 2, L07		L0774: 2, L08	L0655: 2, L08	1,0664: 2, 1,06	S0428: 2, H0725: 2,	H0519: 2, S0126: 2,	H0690: 2, H0	H0684: 2, S0328: 2,	H0522: 2, H0696: 2,	S0390: 2, S0028: 2,	L0748: 2, LO	L0731: 2, L07	H0444: 2, S0436: 2,	10592: 2, 1.0	S0011: 2, H0136: 2,	H0506: 2, H0	H0170: 1, H0686: 1	S0134: 1, \$0430: 1	H0583: 1, H0650: 1	H0656: 1, S0282: 1,	H0255: 1, H0661:	H0663: 1, H0662:	H0241: 1, H	H0125: 1, S0420:	H0637: 1, H0580:	S0007: 1, S0046: 1,	S0132: 1, S0278: 1
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H0431: 1, H0415: 1,		H0586: 1, H0574: 1,	L0622: 1, T0040: 1,	H0250: 1, H0156: 1,	L0021: 1, H0002: 1,	T0082: 1, T0048: 1,	H0318: 1, H0310: 1,	H0251: 1, H0263: 1,	H0597: 1, L0738: 1,	H0546: 1, H0569: 1,	H0050: 1, H0024: 1,	H0578: 1, H0014: 1,	H0015: 1, L0163: 1,	H0051: 1, S0388: 1,	H0083: 1, H0399: 1,	S6028: 1, H0398: 1,	H0428: 1, L0483: 1,	H0181: 1, H0606: 1,	H0166: 1, H0169: 1,	H0708: 1, H0163: 1,	T0067: 1, H0264: 1,	H0488: 1, H0413: 1,	L0564: 1, H0022: 1,	F0041: 1, H0512: 1,	L0475: 1, H0561: 1,	S0294: 1, L0065: 1,	H0641: 1, H0633: 1,	S0472: 1, S0422: 1,
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H0529: 1, L0536: 1, L0770: 1, L0761: 1, L0771: 1, L0648: 1, L0662: 1, L0803: 1, L0606: 1, L0629: 1, L0636: 1, L0517: 1, L0526: 1, L0517: 1, L0526: 1, L0544: 1, L0384: 1, L0544: 1, L0387: 1, H0697: 1, H0711: 1, H0689: 1, H0711: 1, H0669: 1, H0711: 1, H0672: 1, H0539: 1, H0710: 1, H0555: 1, L0740: 1, L0774: 1, L0740: 1, L077: 1, L0758: 1, L0734: 1, L0758: 1, L077: 1, L0758: 1, L077: 1, L0594: 1, H0668: 1, S0026: 1, H0668: 1, S0026: 1, H0668: 1, S0026: 1,		H0694: 48 L0777: 14.	H0521: 11, L0794: 10, L0748: 10, L0747: 10, S0214: 9, S0360: 8,
	3999	4000	4001
	3-512	3-272	3 - 28/
	HPDOP20R 1824		HPDOU54R 1826
	HPDOP20		HPDOU54

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S0003: 8, H0644: 8,	L0439: 8, H0553: 7,	L0662: 7, L0659: 7,	L0588: 7, H0543: 7,	H0038: 6, L0646: 6,	H0436: 6, L0740: 6,	H0341: 5, S0358: 5,	H0580: 5, H0545: 5,	H0494: 5, L0666: 5,	L0665: 5, L0779: 5,	L0599: 5, H0542: 5,	H0656: 4, H0486: 4,	H0046: 4, H0014: 4,	H0551: 4, S0440: 4,	H0529: 4, L0764: 4,	L0809: 4, S0126: 4,	S0406: 4, L0757: 4,	S0436: 4, L0591: 4,	L0485: 4, L0362: 4,	H0423: 4, H0624: 3,	S0376: 3, S0444: 3,	S0408: 3, S0410: 3,	H0581: 3, H0031: 3,	H0090: 3, S0142: 3,	L0770: 3, L0637: 3,	L0772: 3, L0521: 3,	L0766: 3, L0774: 3,	L0663: 3, L0664: 3,	r0068: 3, H0547: 3,	H0539: 3, L0758: 3,
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H0445: 3, S0194: 3,	H0171: 2, H0556: 2,	S0134: 2, H0650: 2,	H0657: 2, S0418: 2,	H0619: 2, H0411: 2,	H0013: 2, S0010: 2,	10544: 2, H0510: 2,	S6028: 2, S0022: 2,	.0483: 2, T0006: 2,	H0591: 2, H0616: 2,	F0067: 2, H0412: 2,	Γ0042: 2, H0560: 2,	H0646: 2, S0422: 2,	L0804: 2, L0775: 2,	L0805: 2, L0783: 2,	L4775: 2, H0670: 2,	H0648: 2, H0710: 2,	H0555: 2, H0478: 2,	S3012: 2, S0027: 2,	L0754: 2, L0749: 2,	L0753: 2, L0759: 2,	S0434: 2, L0592: 2,	L0604: 2, S0011: 2,	S0276: 2, S0040: 1,	H0716: 1, S0114: 1,	L0760: 1, S0116: 1,	H0662: 1, H0459: 1,	S0442: 1, S0045: 1,	S0476: 1, H0393: 1,	S0278: 1, H0550: 1,
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S0222: 1, H0431: 1,	H0455: 1, H0600: 1,	H0592: 1, H0586: 1,	H0331: 1, H0574: 1,	H0485: 1, T0040: 1,	F0109: 1, H0427: 1,	H0156: 1, H0575: 1,	H0004: 1, H0274: 1,	S0346: 1, H0310: 1,	H0052: 1, L0040: 1,	H0597: 1, H0572: 1,	H0123: 1, H0050: 1,	.0471: 1, H0375: 1,	H0594: 1, H0687: 1,	H0252: 1, H0039: 1,	_0142: 1, H0628: 1,	H0617: 1, H0032: 1,		L0455: 1, H0068: 1,	H0598: 1, H0135: 1,	H0163: 1, H0634: 1,	H0264: 1, H0272: 1,		C0041: 1, H0625: 1,	H0561: 1, S0450: 1,	38: 1, H0509: 1,	H0131: 1, H0654: 1,	44: 1, S0210: 1,	S0002: 1, S0426: 1,	.0451: 1, L0520: 1,
202	H04	H05	H03	H04	. 101	H01	H00	803	H00	SOH HOS	HOI	50.7	HOS	H02	101	90H	HOI	<u>각</u>	HOS	HOI	H02	HO4	00T	SOH	804	H01	S0344:	008	2 8
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L0762: 1, L0763: 1, L0761: 1, L0773: 1, L0768: 1, L0386: 1, L0650: 1, L0375: 1, L0651: 1, L0654: 1, L0776: 1, L0527: 1, L0776: 1, L0527: 1, L0776: 1, L0545: 1, L0787: 1, L4501: 1, S0428: 1, S0216: 1, H0144: 1, H0691: 1, S0148: 1, H0519: 1, H0660: 1, H0666: 1, S0330: 1, S0152: 1, S0330: 1, S0152: 1, L0744: 1, L0750: 1, L0756: 1, L0750: 1, L0756: 1, L0751: 1, H0707: 1, L0594: 1, L0756: 1, L0751: 1, H0707: 1, L0594: 1, L0756: 1, S0196: 1, H0422: 1, S0458: 1 and H0506: 1.				
			Pro-6 to Gly-35, Arg-41 to Glu-47.	
	4002	4003	4004	4005
	1 - 504	2 - 178	294 - 485	94 - 291
	1827	1828	1829	1830
	HPDOZ43R	HPDPC65RA	HPDPC90 HPDPC90RA	HPDPH14R
·	HPDO743	HPDPC65	HPDPC90	HPDPH14

	L0784: 2, H0428: 1, H0644: 1, S0366: 1, L0659: 1, L0636: 1, L0789: 1 and H0658: 1.	L0766: 2 and H0658: 2.		L0794: 4, H0351: 3, H0083: 2, H0520: 2, H0658: 2, H0657: 1, H0341: 1, S0358: 1, L0586: 1, L0021: 1, H0040: 1, H0041: 1, H0641: 1, H0641: 1, L0800: 1, L0768: 1, L0803: 1, L0776: 1, L0803: 1, L0777: 1, H0423: 1, H0422: 1 and S0424: 1.	
			Gln-15 to Arg-20, Arg-35 to Ser-40.	Ile-5 to Lys-20.	Asp-28 to His-35, Phe-43 to Trp-48, Asn-81 to Arg-86.
4006	4007	4008	4009	4010	4011
1 - 420	365 - 3	260 - 463	369 - 581	244 - 456	80 - 463
1831	1832	1833	1834	1835	1836
HPDPK37R	HPDPN83R	нрорр69к	HPDPQ10R	нророл би	HPDPR73R
HPDPK37	HPDPN83	НРОРР69	HPDPQ10	HPDPQ16	HPDPR73

L0770: 4, L0771: 4, L0769: 3, L0757: 3, L0766: 2, L0779: 2, L0758: 2, H0170: 1, H0686: 1, S0476: 1, H0586: 1, L0637: 1, L0774: 1, L0776: 1, L0774: 1, L0776: 1, L0791: 1, H0658: 1, H0696: 1, S3012: 1, S0390: 1, L0747: 1, L0752: 1, L0753: 1, L0759: 1, S0434: 1, L0759: 1, and H0543: 1.	0.047011	L0439: 4, H0658: 2, S0474: 1, L0794: 1, L0804: 1, L0791: 1,	L0755: 1 and L0599: 1.	, in the second					
		Asp-17 to Asn-30.				Lys-32 to Glu-39, Glu-43 to Ala-50, Ala-52 to Glu-80.			
4012	4013	4014		4015	4016	4017	4018	4019	4020
2 - 214	2 - 421	378 - 533		2 - 262	2 - 418	148 - 405	1 - 693	2 - 235	2 - 334
1837	1838	1839		1840	1841	1842	1843	-	4-4
HPDPS51R	HPDPS90R	HPDPT86R		HPDPU66R		HPDPY83 HPDPY83RP 00B	TrpDCC34D	REDOCATE NO.	
HPDPS51	06SdCdH	HPDPT86		99ПДОДН	HPDPX12	HPDPY83	7 000 000	HPDOC34	HPDQH11

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	H0052: 11, L0439: 10,	L0748; 7, L0438; 6,	L0750: 6, L0666: 5,	L0599: 5, S0358: 4,	H0620: 4, H0038: 4,	H0413: 4, L0369: 4,	L0769: 4, L0659: 4,	L0665: 4, S0328: 4,	L0758: 4, S0418: 3,	S0442: 3, S0410: 3,	H0619: 3, H0599: 3,	L0662: 3, L0768: 3,	L0774: 3, L0653: 3,	H0660: 3, H0521: 3,	L0731: 3, S0434: 3,	L0601: 3, S0026: 3,	S0424: 3, H0556: 2,	T0002: 2, H0295: 2,	H0583: 2, H0656: 2,	S0376: 2, S0444: 2,	H0580: 2, S0045: 2,	S0046: 2, S0222: 2,	H0592: 2, H0586: 2,	N0009: 2, H0013: 2,	H0575: 2, H0618: 2,	H0318: 2, H0196: 2,	H0012: 2, H0373: 2,
Gln-12 to Ser-19, Gln-40 to Gln-54.								-																			·
4021	4022	•												-													
105 - 308	3 - 545																			•				-			
1846	1847																•										
НРDQH34 НРDQH34R	HPDQI50R													•													
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F0010: 2, S6028: 2,	S0214: 2, H0031: 2,	.0456: 2, H0708: 2,	H0551: 2, H0264: 2,	S0440: 2, H0509: 2,	10529: 2, L0763: 2,	.0638: 2, L0637: 2,	.0764: 2, L0381: 2,	783: 2,	,0663: 2, S0152: 2,	H0478: 2, S0028: 2,	,0741: 2, L0777: 2,	L0588: 2,	605: 2	.0604: 2, H0665: 2	H0136: 2, H0543: 2	H0170: 1, H0265: 1	0717:1	0417: 1	0381: 1	0341: 1	0405:	0356: 1	0208: 1	0411: 1	[0613:	10497:	H0333: 1, H0643:	H0574: 1, L0623: 1,	r0039: 1, H0427: 1,
: 2, S6	: 2, H0	: 2, HO	1: 2, HC	: 2, H0	2, TO	: 2, LO	: 2, L0	7: 2, LO	3: 2, 50	8: 2, SC	1: 2, LC	.0596: 2, LC	.0589: 2, LC	4: 2, H(6: 2, H	0: 1, H	S0040: 1, H0717:	10657: 1, L0417:	J0785: 1, H0381:	S0116: 1, H0341:	H0664: 1, H0402:	H0450: 1, S0356:	S0360: 1, H0208: 1	6: 1, H	H0409: 1, H0613:	HO587: 1, H0497:	13: 1, E	74: 1, I	9: 1, H
T0010	S0214	10456	H0551	S0440	H0529	10638	1,0764	L0657	10663	H047	1074	1.059	1.0589	1060	H013	H017	S004	H065	1078	S011	H066	H045	S036	S047	H040	H058	H033	HOS	1002
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L0021: 1, T0082: 1,	H0036: 1, H0253: 1,	S0010: 1, T0048: 1,	S0182: 1, S0474: 1,	H0263: 1, T0110: 1,	L0040: 1, H0327: 1,	L0471: 1, H0010: 1,	H0024: 1, H0015: 1,	H0416: 1, H0290: 1,	H0292: 1, S0003: 1,	L0483: 1, H0030: 1,	H0553: 1, H0674: 1,	S0364: 1, S0366: 1,	H0059: 1, S0038: 1,	L0564: 1, T0042: 1,	H0494: 1, S0016: 1,	•	H0646: 1, S0426: 1,	•	•	L0772: 1, L0372: 1,	•	•	•	•	L0511: 1, L0656: 1,		•	 H0520: 1, H0547: 1,
																												
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10711: 1,	, H0658: 1, l, H0648: 1,	0378: 1,	10518: 1,	30406: 1,	1018/: 1,	0032: 1,	0.075	.0780: 1,	.0759: 1,	30436: 1,	.0581: 1,	L0593: 1, $ $.0366: 1,	H0542: 1,	H0422: 1 and H0352: 1.	L0779: 5, L0803: 4,	L0663: 3,	J0777: 3, H0687: 2,	H0328: 2, H0038: 2,	,0521: 2, L0809: 2,	L0589: 2,	S0046: 1,	L0717: 1,	H0550: 1,	H0586: 1,	H0574: 1, H0042: 1,	H0544: 1,	Н0620: 1, Н0014: 1,
	H0659: 1, H0658: H0666: 1, H0648:	H0672: 1, S0378:	L0602: 1, H0518: 1,	H0522: 1, S	H0555: 1, H018/:	5002/: 1, 50032:	1,1	L0747: 1, L0780: 1	L0757: 1, I	S0031: 1, S0436:	L0592: 1, L0581:	L0608: 1, 1		H0667: 1,	H0422: 1 8	L0779: 5	H0170: 3, L0663: 3,	L0777: 3,	H0328: 2,	10521: 2,	L0759: 2, L0589: 2	H0171: 1, S0046: 1	S0300: 1, L0717: 1	H0369: 1, H0550:	S0222: 1, H0586: 1	H0574: 1,	S0010; 1, H0544; 1	H0620: 1,
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H0373: 1, H0510: 1, S0003: 1, H0644: 1, H0316: 1, S0036: 1, H0569: 1, L0762: 1, L0662: 1, L0794: 1, L0649: 1, L0659: 1, L0718: 1, L0783: 1, H0547: 1, H0649: 1, H058: 1, L0747: 1, L0780: 1, L0731: 1, L0780: 1, L0731: 1, L0588: 1, L0591: 1,		L0774: 3, L0809: 2, H0483: 1, H0306: 1, H0599: 1, H0618: 1, H0617: 1, H0644: 1, H0617: 1, L0369: 1, L0763: 1, L0743: 1, L0662: 1, L0794: 1, L0665: 1, H0658: 1, H0696: 1, L0748: 1, L0780: 1 and L0757: 1.	L0777: 21, L0748: 16, L0439: 12, S0360: 9,
		Pro-5 to Arg-25, Lys-32 to Ser-39, Pro-63 to Pro-79, Pro-115 to His-121.	
	4024	4025	4026
	2-646	256 - 711	1 - 639
	1849		1851
	HPDQI70R	HPDQL55R	HPDQR20 HPDQR20R
	HPDQI70	HPDQL55	HPDQR20

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0659: 9, L0666: 9,	0803: 8, L0662: 7,	747: 7, L0794: 6,	.0663: 5, S0406: 5,	770: 4, L0731: 4,	0758: 4, S0026: 4,	S0356: 3, L0717: 3,	10156: 3, L0769: 3,	0809: 3, L0664: 3,	10520: 3, H0672: 3,	H0521: 3, H0696: 3,	.0750: 3, S0192: 3,	H0542: 3, H0423: 3,	H0624: 2, H0657: 2,	S0420: 2, S0358: 2,	S0376: 2, S0408: 2,	H0013: 2, S0003: 2,	S0214: 2, H0264: 2,	H0412: 2, H0413: 2,	S0440: 2, L0764: 2,	.0766: 2, L0804: 2,	774: 2, L0775: 2,	0806: 2, L0655: 2,	0665: 2, S0374: 2,	.0438: 2, H0547: 2,	10519: 2, H0689: 2,	H0660: 2, S0380: 2,	H0522: 2, S0206: 2,	.0749: 2, L.0752: 2,	.0759: 2, L0361: 2,
07	<u> </u>	107	<u> </u>	5		203	OH	<u> </u>	HO	HO	2	OH	H	<u>80</u>	SOS	HO	SO.	<u> </u>	SO	<u>2</u>	<u>2</u>	<u>3</u>	ន	<u>3</u>	OH OH	<u>H</u>	OH HO	3	C.
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S0242: 2. H0170: 1.	H0171: 1, L0448: 1	H0685: 1, H0716: 1	S0114: 1, H0656: 1,	.0785: 1, S0400: 1,	S0442: 1, S0444: 1,	, S0222: 1,	, H0586: 1	H0632: 1, L0623: 1,	, H0036: 1	H0581: 1, H0309: 1	, H0327: 1	H0545: 1, H0150: 1	, H0024: 1,	H0014: 1, T0010: 1,	, H0252: 1	, H0688: 1	H0622: 1, L0483: 1,	, S0366: 1,	, H0163: 1	H0090: 1, H0040: 1	, H0623: 1,	r0069: 1, H0494: 1,	S0448: 1,	H0714: 1, H0641: 1	S0142: 1, S0210: 1,	, L0646: 1,	L0771: 1,	.0387: 1, L0651: 1,	, L0653: 1,
S0242: 2	H0171: 1	H0685: 1	S0114: 1,	L0785: 1	S0442: 1,	S0300: 1,	H0431: 1	H0632: 1	H0069: 1	H0581: 1	H0263: 1	H0545: 1	1.0471: 1	H0014: 1	H0083: 1	H0615: 1	H0622: 1	H0068: 1	H0598: 1	H0090: 1	Т0067: 1	T0069: 1,	S0372: 1,	H0714: 1	S0142: 1,	L0369: 1,	1,0641: 1,	L0387: 1,	L0378: 1,
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L0542: 1, L0647: 1, L0787: 1, H0658: 1, H0670: 1, H0648: 1, H0539: 1, H0710: 1, S0332: 1, H0478: 1, H0627: 1, L0755: 1, S0436: 1, L0605: 1, L0362: 1, H0667: 1, S0276: 1, H0543: 1, H0422: 1, H0506: 1 and					H0658: 2, L0745: 2, S0422: 1, H0648: 1 and L0731: 1.	L0748: 9, L0774: 3, H0331: 2, H0632: 2, L0526: 2, H0658: 2, L0581: 2, H0574: 1 and L0790: 1.	H0542: 11, H0265: 7, H0543: 7, L0751: 5, S0424: 5, H0556: 4,
	Gln-121 to Glu-136, Ala-153 to Ser-162, Ala-176 to Ser-182.			Phe-25 to Arg-34, Cys-51 to His-56, Glu-66 to Ser-75.		His-2 to Trp-13, Lys-22 to Cys-31.	Gly-17 to Glu-26, Lys-32 to Leu-37, Phe-46 to Thr-52,
	4027	4028	4029	4030	4031	4032	4033
	2 - 550	113 - 325	25 - 273	25 - 258	134 - 322	41 - 175	3 - 410
	1852	1853	1854	1855	1856	1857	1858
	HPDQR88R	HPDOS14R	HPDOS25R	HPDQS63R	HPDQS66R	HPDQS76R	HPDQT32R
	HPDQR88	HPDOS14	HPDOS25	HPDQS63	нРDQS66	HPDQS76	HPDQT32

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S0420: 4, S0010: 4, S0143: 4	0743: 4, L0747: 4,	.0753: 4, H0657: 3,	10370: 3, H0618: 3,	10251: 3, H0545: 3,	10457: 3, H0620: 3,	10644: 3, H0560: 3,	.0775: 3, L0663: 3,	10672: 3, S3012: 3,	.0748: 3, S0212: 2,	S0418: 2, S0360: 2,	10580: 2, S0222: 2,	H0427: 2, H0253: 2,	S0049: 2, H0327: 2,	.0471: 2, H0617: 2,	10124: 2, S0366: 2,	10412: 2, L0667: 2,	.0655: 2, H0690: 2,	10682: 2, H0658: 2,	H0670: 2, H0660: 2,	H0627: 2, S0027: 2,	.0741: 2, L.0740: 2,	H0506: 2, H0170: 1,	.0002: 1, H0686: 1,	H0381: 1, H0484: 1,	H0255: 1, H0663: 1,	H0664: 1, H0638: 1,	S0358: 1, H0675: 1,	H0637: 1, S0468: 1,
109, S0420	וונ	1.075	H037	H025	H045	H064	1,077	190H	1074	S0418	H058	H042	S004	1047	H012	H041	1065	890H	H067	H062	L074	H050	T000(H038	H025	990H	S035	H063
Ile-103 to Gly-109,	Lys-120 to Om																											
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, S0132: S0005: , H0156: , H0318: , T0103:)1 /8)051:)355)594)687)194	0006 0606 0674	0063 0067 0494 0352 0633	1, S0144: 1, S0426: 1, L0369: 1, L0646: 1, L0662: 1, L06	L0376: 1 L057: 1 L0542: 1 L0787: 1
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H0208: 1, S0132: 1 S0278: 1, S0005: 1 H0497: 1, H0156: 1 L0021: 1, H0318: 1 H0052: 1, T0103: 1	10544; 1, h0178; 1 10172; 1, 80051; 1 10083; 1, H0355; 1 H0510; 1, H0594; H0188; 1, H0687; H0687; 1	20483: 1, T0006: 1 H0033: 1, H0424: H0031: 1, H0606: H0166: 1, H0674:	H0591: 1, H0063: 1 H0087: 1, T0067: 1, L0351: 1, H0494: 1, H0561: 1, S0352: 1, S0440: 1, H0633: 1,	H0647: 1, S0144: 1, S0344: 1, S0426: 1, H0529: 1, L0369: 1, L0372: 1, L0646: 1, L0800: 1, L0662: 1, L058: 1, L0568: 1, L058: 1, L	
201 201 201 201 201 201 201 201 201 201	H0544: 1, H0178: 1, H0172: 1, S0051: 1, H0083: 1, H0355: 1, H0510: 1, H0594: 1, H0188: 1, H0687: 1, H0622: 1, L0194: 1,		S H L S H S	222232 222232	3333
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L0666: 1, L0664: 1, L0665: 1, H0593: 1, H0684: 1, H0435: 1, H0659: 1, S0328: 1, S0330: 1, H0539: 1, S0378: 1, H0631: 1, H0555: 1, H0631: 1, L0774: 1, L0750: 1, L0777: 1, L0752: 1, L0757: 1, L0758: 1, L0596: 1, L0485: 1, L0593: 1, L0601: 1, S0011: 1, S0026: 1, S0192: 1 and H0352: 1.	L0731: 4, L0775: 3, L0740: 3, H0170: 2, S0426: 2, L0770: 2, L0646: 2, L0766: 2, L0754: 2, H0344: 1, H0592: 1, H0393: 1, H0592: 1, H0318: 1, S6028: 1, H0038: 1, L0776: 1, L0642: 1, L0776: 1, L0663: 1, L0776: 1, L0774: 1, L0778: 1, L0774: 1, L0778: 1, L0777: 1, L0748: 1, L0777: 1, L0752: 1, L0777: 1, H0645: 1, S0424: 1 and S0384: 1.
	Pro-25 to Thr-32.
	4034
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	1859
	HPDQU81R
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		AR104: 42, AR089: 22, AR060: 7, AR096: 4, AR055: 2, AR061:															H0031: 2, H0135: 1,	DOZIZ. I AMA INOSO: 1:				
		Asn-20 to Asp-25, AR104: 42, AR089: Met-75 to Asp-81, 22, AR060: 7, AR096 A 75-144 to Thr-152 4, AR055: 2, AR061:	Lys-1 to Leu-23, Ala-27 to Ser-34,	Gly-36 to Phe-44.				Leu-11 to Cys-24,	Leu-30 to Ser-36,	Gly-93 to Glu-99.		Ser-120 to Tyr-126,	Gly-137 to Arg-142, Ser-160 to Ser-168.			Glu-1 to Pro-9.			Ala-2 to Gly-8.			Gly-1 to Cys-24,
4035	4036	4037	4038		4039	4040	4041	4042			4043	4044		4045	4046	4047	4048		4049	4050	4051	4052
2-421	246 - 401	68 - 550	113 - 391		3 - 476	3 - 437	164 - 340	29 - 430			212 - 538	3 - 506		1.505	1 - 384	10			2 - 481	1 - 177	3 - 302	2 - 307
1860	1861	1862	1863		1864	1865	1866	1867			1868	1869		1870	1871	1872	1873		1874	1875	1876	
atomoran			HPDQX13R		HPDQY23R	HPDQY65R	HPDOY88R	HPDOY95R	,		HPDOZ30R	HPDOZ65R	,	TIPLE A 44B	Ans A gridu	HPDRE65R	HPDRG34R		HPDRG73R	HPDRM7	HPDRM9	HPDRO04
-	HPDOV0/	HPDQW39	нРDQX13		HPDOY23	HPDOY65	HPDOY88	HPDOY95			HPDO730	HPDOZES	2007	77.7	HPDRA#	אפתחמט	HPDRG34		HPDRG73		HPDRM93	HPDRO04

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				S0132: 1, H0156: 1,	H0575: 1, H0551: 1,	H0658: 1, H0670: 1,	H0555: 1 and L0439: 1.	L0766: 11, L0752: 7,	L0754: 5, H0547: 4,	50358: 3, 50022: 3,	L0756: 3, L0777: 3,	H0625: 2, L0065: 2,	S0438: 2, L0805: 2,	L0776: 2, H0658: 2,	S0328: 2, L0740: 2,	L0745: 2, L0747: 2,	L0750: 2, L0731: 2,	L0601: 2, H0542: 2,	H0543: 2, H0423: 2,	S0424: 2, H0341: 1,	H0661: 1, H0664: 1,	H0306: 1, H0402: 1,	H0638: 1, S0360: 1,	H0580: 1, H0645: 1,	H0443: 1, H0357: 1,	H0592: 1, H0587: 1,	H0574: 1, H0486: 1,	H0270: 1, H0156: 1,	T0048: 1, H0052: 1,
Cys-27 to Gly-43,	Ala-46 to Trp-54,	Ala-56 to Arg-68,	Phe-83 to Arg-93.	Thr-4 to His-17,	Ser-21 to Gln-27.	-		Thr-57 to Phe-62,	Gly-68 to Phe-73,	His-86 to Tyr-92,	Asp-97 to Phe-103.		-																
		٠		4053				4054																					
		:		249 - 509				3 - 323		*		,									•		·						
				1878				1879																					
	•			HPDRO64R				HPDRP36R							•														
				HPDR064				HPDRP36																	_				

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H0375: 1, H0687: 1, H0328: 1, H0169: 1, H0598: 1, H0551: 1, H05413: 1, T0042: 1, H0559: 1, L0520: 1, L0536: 1, L0772: 1, L0649: 1, L0666: 1, L0665: 1, H0144: 1, H0520: 1, H0519: 1, H0559: 1, H0651: 1, S0380: 1, H0696: 1, S0380: 1, H0696: 1, L0749: 1, L0753: 1, L0749: 1, L0753: 1, L0757: 1, L0759: 1, H0445: 1, S0436: 1, S0436: 1, L0594: 1 and H0422: 1.		L0748: 3, H0556: 2, S0420: 2, S0045: 2, H0171: 1, H0265: 1, S0134: 1, H0657: 1,
	Gly-8 to Val-14, Gln-20 to Asp-27, Met-79 to His-88, Arg-171 to Tyr-183, Pro-198 to Gly-204.	
	4055	4056
	2 - 622	27 - 290
	1880	1881
	HPDRP41R	HPDRQ66R
	HPDRP41	HPDRQ66

H0341: 1, S0110: 1, H0661: 1, H0305: 1, H0580: 1, S0007: 1, S0046: 1, S0132: 1, S0278: 1, H0257: 1, H0069: 1, H0575: 1, H0252: 1, H0634: 1, H0412: 1, H0634: 1, H0633: 1, S0344: 1, H0639: 1, L0521: 1, L0806: 1, L0805: 1, H0435: 1, H0658: 1, H0672: 1, L0599: 1, L0747: 1, L0599: 1,						
	Gly-6 to Ser-16, Glu-19 to Ala-33, Glu-44 to Lys-60, Ile-135 to Lys-147.	Ala-3 to Lys-11, Gly-112 to Glu-117.	Lvs-94 to Ser-100.	Lys-32 to Glu-38, Ser-44 to Ser-55,	Gln-67 to Gly-78, Glu-85 to Glu-90,	Gly-108 to Ser-114, Phe-149 to Lys-158.
	4057	4058	4059	4061	,	
	1 - 441	1 - 543	3 - 332	2 - 508		
	1882	1883	1884	1886		
	HPDRQ84R	HPDRR71R	HPDRS43R HPDRS46R	1		
	HPDRQ84	HPDRR71	HPDRS43	HPDRS87		

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L0754: 19, L0731: 18, H0521: 15, L0777: 11, L0766: 8, L0755: 7,	H0581: 6, L0659: 6, L0751: 6, H0486: 5,	H0266: 5, H0553: 5,	H0144: 5, L0439: 5,	S0276: 5, H0638: 4,	S0360: 4, H0644: 4,	S0374: 4, S0330: 4,	.0740: 4, L.0756: 4,)599: 4, L0362: 4,	H0423: 4, H0657: 3,	H0663: 3, H0580: 3,	H0575: 3, S0003: 3,	H0622: 3, H0124: 3,	H0264: 3, H0623: 3,	S0002: 3, L0520: 3,	0764: 3, L0775: 3,	.0655: 3, L0519: 3,	10519: 3, H0659: 3,	10658: 3, S0328: 3,	.0744: 3, L0747: 3,	,0759: 3, S0026: 3,	S0194: 3, H0624: 2,	H0656: 2, H0341: 2,	S0354: 2, S0408: 2,	S0046: 2, H0619: 2,	H0013: 2, H0250: 2,
4062 Lys-25 to Arg-36. L HC LO	<u>H</u> 01	H	<u>i</u> ii	S	SC	SC	ĭ	<u>a</u>		H	H	<u> </u>	正	S	<u>.</u>	<u></u>	<u> </u>	<u>. 11</u>		<u>•</u> 1	<u> </u>			<u> </u>	
4062																									
59 - 214																									
1887																									
HPDRT36R			•				-	•																	
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H0545: 2, L0471: 2,	H0428: 2, L0142: 2,	H0551: 2, T0067: 2,	H0561: 2, H0641: 2,	30210: 2, L0772: 2,	.0771: 2, L0794: 2,	J0803: 2, L0661: 2,	JO544: 2, L0793: 2,	.0666: 2, L.0665: 2,	S0216: 2, L0565: 2,	10648: 2, S0152: 2,	10522: 2, H0555: 2,	10345: 2, S0027: 2,	.0750: 2, L0779: 2,	.0752: 2, L.0758: 2,	10445: 2, S0434: 2,	.0591: 2, L0594: 2,	10653: 2, H0422: 2,	H0159: 1, S0134: 1,	H0583: 1, H0650: 1,	S0212: 1, H0661: 1,	H0662: 1, H0402: 1,	S0358: 1, S0444: 1,	T0008: 1, S0045: 1,	S0132: 1, H0411: 1,	H0369: 1, H0370: 1,	70039: 1, T0060: 1,	H0042: 1, H0590: 1,	0048: 1, H0318: 1,	H0596: 1, L0738: 1,
4	<u> </u>	<u></u>	<u>144</u>	<u>01</u>	H		<u>1</u>		<u> </u>	<u> </u>	<u>, 11</u>	<u>.</u> ±4	<u>. </u>		<u> </u>		<u> 144.</u>	<u>,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,</u>		S	<u> </u>	<u>v</u>	_=_	<u>S</u>	<u> </u>	E	<u>.</u>	<u> Frai</u>	14
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.0157: 1,	10024: 1,	10267: 1,	0214: 1,	H0030: 1,	H0673: 1,	H0316: 1,	H0090: 1,	H0040: 1,	H0488: 1,	H0412: 1,	H0056: 1,	H0494: 1,	10646: 1,	S0344 <u>:</u> 1,	.0371: 1,	L0800: 1,	L0648: 1,	L0389: 1,	L0774: 1,	L0653: 1,	L0658: 1,	L0809: 1,	H0701: 1,	S0126: 1,	H0683: 1,	, S0454: 1,	H0479: 1,	L0743: 1,	H0343: 1,
H0046: 1, L0157: 1	H0565: 1, H0024:	S0388: 1, H0267: 1	80250: 1, 80214: 1,	H0328: 1, 1	H0383: 1, 1	H0169: 1, 1	H0163: 1, 1	H0591: 1, 1	H0634: 1, 1	H0268: 1, 1	H0413: 1, H0056:	L0435: 1, H0494: 1	S0370: 1, I	H0652: 1,	S0426: 1, L0371:	L0646: 1, 1		L0768: 1,]	L0552: 1,	L0805: 1, L0653: 1,	L0776: 1,	1.0542: 1,	1, 1, 1, 1,	H0691: 1,	H0711: 1, H0683:	H0518: 1,	S0013: 1, H0479: 1,	S0206: 1,	S0260: 1, H0343: 1,
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L0584: 1, H0542: 1 and	80412: 1.					-								L0794: 6, L0758: 6,	L0747: 5. L0770: 4.	L0809: 4, H0620: 3,	L0764: 3, H0658: 3,	H0618: 2, H0012: 2,	H0135: 2, L0769: 2,	L0771: 2, L0773: 2,	L0662: 2, L0803: 2,	L0532: 2, S0126: 2,	L0756: 2, S6024: 1,	S0282: 1, H0638: 1,	S0420: 1, L0717: 1,	S0222: 1, H0441: 1,	H0333: 1, L0622: 1,	H0309: 1, H0597: 1,	S0388: 1, S0051: 1,
		Ala-2 to Asp-9,	Arg-18 to Gly-26,	Asn-34 to Arg-43,.	Ala-52 to Asp-57.	Leu-48 to Lys-63.		Glu-11 to Pro-16,	Arg-46 to Tyr-56,	Asp-63 to Asp-69.		Glu-54 to Val-64,	Leu-97 to Leu-105.	Ser-14 to Gly-20.						,								•	
		4063				4064	4065	4066			4067	4068		4069										•			-		
		3 - 560				2 - 196	2 - 334	1 - 297			458 - 703	1 - 528		965 - 008		٠					_	•							
		1888				1889	1890	1891			1892	1893		1894															
		HPDRT37R				HPDRT38R	HPDRU03R	HPDRU37R			HPDRU71R	HPDRV73R		HPDRV79R			-												
		HPDRT37				HPDRT38	HPDRU03	HPDRU37			HPDRU71	HPDRV73		HPDRV79				٠						•					

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T0010: 1, H0271: 1, H0687: 1, L0483: 1, H0604: 1, H0630: 1, H0412: 1, L0667: 1, L0646: 1, L0800: 1, L0649: 1, L0374: 1, L0649: 1, L0374: 1, L0652: 1, L0804: 1, L0652: 1, L0607: 1, L0659: 1, L0559: 1, L0659: 1, L0577: 1, L0666: 1, L0663: 1, L0666: 1, L0663: 1, L0666: 1, L0789: 1, L0666: 1, L0789: 1, L0665: 1, H0670: 1, H0627: 1, L0781: 1, L0677: 1, L0731: 1, L0605: 1, H0670: 1, H0657: 1, H0423: 1, L0605: 1, H0423: 1, L0605: 1, H0423: 1, L0697: 1 and S0462: 1.			
	4070 Ala-31 to Gln-37, Lys-63 to Ser-69.		Pro-3 to Gly-8, Pro-70 to Pro-85.
·	4070	4071	4072
	1 - 231	192 - 587	2 - 322
	1895	1896	1897
	HPDRW09R	HPDRW51R	HPDRY37
	HPDRW09	HPDRW51	HPDRY37

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	H0457: 9, L0766: 7,	H0046: 5, H0650: 4,	H0634: 3, L0659: 3,	L0439: 3, H0716: 2,	H0656: 2, H0254: 2,	H0255: 2, H0333: 2,	H0253: 2, H0081: 2,	H0083: 2, H0628: 2,	L0761: 2, L0800: 2,	H0658: 2, L0750: 2,	H0543: 2, H0422: 2,	H0265: 1, H0657: 1,	S0001: 1, S0420: 1,	S0354: 1, S0376: 1,	S0360: 1, H0722: 1,	S0045: 1, H0550: 1,	S0222: 1, H0614: 1,	H0613: 1, H0069: 1,	H0427: 1, H0575: 1,	H0318: 1, H0581: 1,	H0052: 1, H0327: 1,	H0545: 1, H0041: 1,	H0050: 1, H0354: 1,	H0266: 1, H0179: 1,	H0428: 1, H0622: 1,	S0366: 1, S0036: 1,	H0087: 1, H0264: 1,	H0488: 1, L0435: 1,
Arg-13 to Gly-26, Asn-35 to Gln-40.	Val-30 to Trp-36,	Ala-53 to Glu-67.										-			3.													
4073	4074								•																			
1 - 354	3 - 509								-																		,	
1898	1899																	-		-								
	HPDRZ29R				-								•															
HPDRZ11	HPDRZ29		-																				-					

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H0022: 1, T0041: 1, H0560: 1, S0142: 1, S0426: 1, H0695: 1, H0529: 1, L0644: 1, L0649: 1, L0662: 1, L0805: 1, L0658: 1, L0805: 1, L0538: 1, L0792: 1, L0539: 1, L0663: 1, L0664: 1, H0572: 1, S0330: 1, H0551: 1, S0406: 1, H0555: 1, H0436: 1, S0027: 1, S0028: 1, L0744: 1, L0749: 1, L0756: 1, H0445: 1, S0434: 1, H0136: 1, S0434: 1, H0136: 1,		T 07/10: 10 T 08/03: 3	LO439: 3, L0751: 3, L0626: 2, L0809: 2, L0790: 2, H0658: 2, L0744: 2, L0740: 2, L0754: 2, L0747: 2,
			Pro-2 to Lys-18.
	4075	4076	4077
	774 - 484	243 - 380	40 - 315
		1901	1902
·	HPDVA01R	1	HPDVB37R
	HPDVA01	HPDVA06	HPDVB37

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L0758: 2, H0170: 1, H0645: 1, H0351: 1, H0370: 1, H0497: 1, S0388: 1, T0067: 1, L0772: 1, L0643: 1, L0804: 1, L0774: 1, L0776: 1, L0666: 1, L0663: 1, L0665: 1, L0438: 1, S0044: 1,	L0745: 1, L0749: 1, L0755: 1, L0757: 1 and S0031: 1.															
				Pro-8 to Pro-13.	Gln-22 to Val-33,	Pro-37 to Gly-45,	Ser-56 to Thr-65,	Arg-82 to Thr-92.	Glu-79 to Gln-86.	Met-32 to Lys-37,	Asp-44 to Ala-60.	Ala-32 to Thr-42,	Lys-55 to Phe-60,	Asp-85 to Gly-90,	Arg-95 to Thr-100.	Ile-20 to Gly-29, Asn-82 to Thr-87.
		4078	4079	4080	4081				4082	4083		4084				4085
		2 - 778	50 - 136	89 - 220	3 - 326				193 - 450	1-510		3 - 338	-			3 - 425
		1903	1904	1905	1906				1907	1908		1909				1910
		HPDVB70R	HPDVC28R	HPDVE55R	HPDVG06 HPDVG06R				HPDVG67R	HPDV125R		HPDVI95R				HPDVK74 HPDVK74R
		HPDVB70	HPDVC28	HPDVE55	HPDVG06				HPDVG67	HPDVI25		HPDV195				HPDVK74

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		10.1	218: 1, 6: 1, 33: 1,	27: 1, 45: 1,	87: 1,	47: 1, 58: 1,	572: 1,	39: 1,	HU4425.	10. 1,	194.	74: 1,	558: 1,	757: 1,	H0543:							
		200	H0656: 3, SUZ18: 1 S0116: 1, S0356: 1, H0586: 1, H0333: 1,	H0486: 1, H0427:	H0135: 1, H0087:	H0695: 1, H0547: 1 H0435: 1, H0658: 1	H0670: 1, H0672: 1,	H0521: 1, L0439: 1	H0543: 1 and H0423: 1	HU656: 1, 50110: 1	503/0. 1, 10110: -1	0649: 1, L0774: 1,	.0665: 1, H0658: 1	J0748: 1, L0757: 1	H0667: 1 and H0543:							
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Arg-8 to Lys-13, Gly-35 to Lys-42, Ala-48 to Lys-54.			Gly-1 to Phe-8, Pro-39 to Thr-61, Pro-66 to Phe-71.	Arg-84 to Pro-89.						Ala-5 to Val-11,	Leu-15 to Lys-23,	Lys-39 to 11e-44.		-		Lys-13 to Leu-24,	Arg-31 to Pro-54.	Ala-34 to Thr-46,	Asp-59 to Lys-73,	Ala-86 to Gin-94,	Pro-99 to Gln-109,	Glu-129 to 1 hr-141
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Asp-160 to Asn- 168.	Ala-33 to Cys-41,	Thr-58 to Pro-66.			•																							
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	Glu-1 to Asp-6, Thr-11 to Glu-20, Val-61 to Gln-66.	Pro-1 to Ala-9, Arg-17 to Glu-25.	1			Val-1 to Glu-9, Pro-26 to Gly-31, Tyr-67 to Glu-74,
	4094	4095	4096	4097	4098	4099
·	1 - 465	3 - 278	29 - 421	3 - 284	1 - 333	91 - 693
	1919	1920	1921	1922	1923	
	HPDVO67 HPDVO67R	HPDVQ74R	HPDVO82R	HPDVT30R	HDDV/T37R	HPDVU28R
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Phe-107 to Glu-118, Lys-162 to Arg-173.	Ser-1 to His-7,	Thr-15 to Arg-29,	His-31 to Val-42,	Pro-52 to Pro-94,	Glu-113 to Ala-131,	Lys-161 to Arg-167. L0665: 2, H0693: 2,	,						<u> </u>					<u> </u>					<u> </u>					
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		-1 -	 1.	1	Pro-6 to Ala-16.				Arg-1 to Glu-8,	Ala-31 to 115-27.		Ala-3 to Ala-10,	Pro-15 to Pro-25,	Glu-79 to Ser-88,	Pro-115 to Arg-122.													
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9: 1,	6: 1,	8:1,	9:1,	5: 1,	1: 1,	5: 1,	7:1,	1: 1,	5: 1,	2: 1,	8. 1.	3: 1,	4.	3: 1,	6: 1,	H0135: 1,	4: 1,	H0087: 1,	4.	1	4: 1	10641: 1,	4.1	6: 1,	.0646: 1, 1	,0648: 1,]	2: 1,	5: 1,	1.1
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L0807: 1, L0542: 1, L0519: 1, L0530: 1, L0790: 1, L0663: 1, H0144: 1, H0724: 1, H0660: 1, H0666: 1, H0521: 1, S0190: 1, H0134: 1, H0187: 1, H0134: 1, H0187: 1, L0779: 1, L0751: 1, S0028: 1, L0751: 1, S0031: 1, H0445: 1, S0436: 1, H0668: 1, S0424: 1.			H0575: 2, L0002: 1, S0336: 1, H0487: 1, L0762: 1, L0637: 1, L0805: 1, L0659: 1, H0658: 1, L0777: 1 and L0731: 1.		H0038: 5, H0616: 3, L0752: 3, S0418: 2, H0620: 2, H0547: 2, L0602: 2, H0521: 2,
	Asn-60 to Gly-72, Pro-83 to Trp-114, Ser-117 to Arg-126.	Arg-1 to Gln-9.		-	Phe-1 to Cys-13, Gly-28 to Ser-35.
	4111	4112	4113	4114	4115
	1 - 378	2 - 121	248 - 559	1 - 507	169 - 342
	1936	1937	1938	1939	
	HPDWD81R	HPDWE11R	HPDWE64	HPDWF93R	HPDWG5
	HPDWD81	HPDWE11	HPDWE64	HPDWF93	HPDWG51

	·	
L0751: 2, S0212: 1, S0282: 1, H0638: 1, S0358: 1, S0358: 1, H00518: 1, H0261: 1, H0275: 1, S0346: 1, H0576: 1, H0071: 1, H0510: 1, S0003: 1, H0549: 1, S0144: 1, H0649: 1, S0144: 1, H0658: 1, H0518: 1, S0013: 1, S0044: 1, L0743: 1, L0743: 1, L0743: 1, L0743: 1, L0743: 1, L0759: 1, L0599: 1 and S0424: 1, L0599: 1 and S042	H0556: 30, H0659: 9, L0439: 9, S0406: 8, H0494: 7, S0360: 6, L0771: 6, H0634: 5, S0206: 5, L0777: 5, H0265: 4, H0657: 4,	S0328: 4, S0152: 4, S0146: 4, L0748: 4, L0751: 4, L0749: 4, H0423: 4, S0040: 3, H0497: 3, H0623: 3,
	Arg-21 to Gly-32, H0556: 30, H0659 Pro-63 to Ser-79, L0439: 9, S0406: 8, Glu-107 to Glu-126, L0771: 6, H0634: 5, Glu-133 to Ser-146, S0206: 5, L0777: 5, Ser-148 to Thr-157, H0265: 4, H0267: 4	
	4118	
	174 - 308	
	1943	
	HPDWL70 HPDWL70R HPDWN46 HPDWN46R	
	HPDWL70 HPDWN46	

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H0529: 3, L0761: 3,	0665: 3, H0144: 3,	H0555: 3, S0026: 3,	H0656: 2, H0341: 2,	S0212: 2, S0418: 2,	S0356: 2, H0675: 2,	S0046: 2, H0411: 2,	H0586: 2, H0581: 2,	H0615; 2, H0553; 2,	H0591: 2, S0440: 2,	S0422: 2, L0772: 2,	L0438: 2, H0711: 2,	H0521: 2, S0404: 2,	S3014: 2, L0743: 2,	L0740: 2, H0170: 1,	H0171: 1, H0159: 1,	H0686: 1, S0342: 1,	H0294: 1, S0114: 1,	T0049: 1, S0116: 1,	H0661: 1, H0458: 1,	S0420: 1, S0376: 1,	S0444: 1, H0437; 1,	S6022: 1, H0550: 1,	H0362: 1, H0642: 1,	H0574: 1, H0559: 1,	S0280: 1, H0590: 1,	H0004: 1, H0363: 1,	H0263: 1, H0596: 1,	H0081: 1, H0014: 1,	H0688: 1, H0628: 1,
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L0055: 1, H0673: 1, H0068: 1, H0551: 1, H0625: 1, S0002: 1, L0369: 1, L0435: 1, L0369: 1, L0637: 1, L0764: 1, L0774: 1, L0775: 1, L0375: 1, L0805: 1, L0806: 1, L0805: 1, L0657: 1, L0792: 1, H0701: 1, S0126: 1, H0519: 1, S0126: 1, H0548: 1, H0670: 1, H0648: 1, L0779: 1, L0747: 1, L0759: 1, H0548: 1, H0667: 1, S0194: 1, H0667: 1, S0194: 1, H0642: 1 and S0424: 1.			L0766: 5, H0551: 3, L0769: 3, L0639: 3, H0327: 2, S0002: 2, L0764: 2, L0659: 2,
		Gly-2 to Pro-9, Ser-24 to Thr-33.	
	4119	4120	4121
·	1 - 246	2 - 607	351 - 704
	1944	1945	1946
	HPDWN65R	HPDWN69 HPDWN69R	HPDWO08R
	HPDWN65 HPDWN65	HPDWN69	HPDWO08 HPDWO08

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L0663: 2, L0777: 2, S0356: 1, S0360: 1, S0410: 1, L0717: 1, H0261: 1, H0550: 1, H0575: 1, T0109: 1, H0575: 1, T0048: 1, H0052: 1, S0051: 1, L0455: 1, H0124: 1, H0135: 1, H0124: 1, L0770: 1, L0763: 1, L0761: 1, L0643: 1, L0562: 1, L0635: 1, L0565: 1, L0658: 1, H0522: 1, L0756: 1, H0522: 1, L0756: 1,			L0770: 12, H0657: 10, H0659: 9, L0776: 8, H0648: 8, L0755: 8, L0774: 7, S0410: 6, H0494: 6, L0769: 6, L0750: 6, S0408: 5, S0440: 5, L0775: 5,
	His-1 to Gly-13. Arg-90 to Lys-95, Pro-121 to Trp-129.		Ala-3 to Glu-8.
	4122	4124	4125
	386 - 568 27 - 455	2 - 358	1 - 342
	1947	1949	<u> </u>
	HPDWO29 HPDWO29R HPDWO50 HPDWO50R	HPDW061R	HPDWO82
·	HPDWO29 HPDWO50	HPDWO61	

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S0376: 4, S0360: 4,	0748: 4, L0777: 4,	0752: 4, L0759: 4,	10341: 3, S0358: 3,	10616: 3, L0766: 3,	10: 3, L0747: 3,	.0731: 3, L0758: 3,	S0026: 3, H0170: 2,	10556: 2, H0685: 2,	10656: 2, S0420: 2,	S0444: 2, T0109: 2,	H0156: 2, H0545: 2,	.0471: 2, H0031: 2,	H0040: 2, H0560: 2,	.0065: 2, S0438: 2,	.0763: 2, L.0772: 2,	.0764: 2, L0518: 2,	J0783: 2, H0144: 2,	S0374: 2, H0520: 2,	H0658: 2, H0670: 2,	.0753: 2, H0445: 2,	S0434: 2, H0665: 2,	S0194: 2, S0134: 1,	S0116: 1, H0663: 1,	.0481: 1, H0638: 1,	H0125: 1, S0442: 1,	H0351: 1, H0369: 1,	H0550: 1, H0431: 1,	H0331: 1, H0069: 1,	S0346: 1, T0048: 1,
8037	107	L075	H037	190H	107	<u> 1</u> 03	S002	H05	H064	S 044	H015	1047	H004	907	1.076	1076	1078	S037	H065	1.075	S043	S019	S011	1048 84	H012	H035	H055	H033	S034
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S0182: 1, S0049: 1, H0563: 1, H0081: 1, H0620: 1, H0014: 1, H0373: 1, H0083: 1, H0778: 1, H0428: 1, L0475: 1, H0434: 1, L0598: 1, H0529: 1, L0598: 1, L0762: 1, L0371: 1, L064: 1, L0371: 1, L0521: 1, L0573: 1, L0521: 1, L0573: 1, L0521: 1, L0555: 1, L0521: 1, L0555: 1, L051: 1, L0555: 1, L051: 1, L0546: 1, L0547: 1, H0660: 1, H0435: 1, H0660: 1, H0651: 1, S0380: 1, S0406: 1, L0749: 1, L0780: 1, H0668: 1, H0653: 1, H0668: 1, H0422: 1 and H0542: 1, H0422: 1 and	H0617: 8, S0404: 8, H0484: 7, H0550: 7,
· · · · · · · · · · · · · · · · · · ·	Asp-15 to Val-20.
	4126
	73 - 321
	1951
	HPDWS61R
	HPDWS61

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.0758: 5, H0606: 4,	H0658: 4, H0696: 4,	S0276: 4, H0295: 3,	S0280: 3, H0150: 3,	H0188: 3, H0163: 3,	0769: 3, S0408: 2,	.0622: 2, L0623: 2,	10060: 2, T0006: 2,	10181: 2, H0673: 2,	10494: 2, L0764: 2,	.0775: 2, L0776: 2,	.0514: 2, L0510: 2,	.0511: 2, L0512: 2,	.0540: 2, L0783: 2,	.0809: 2, L0528: 2,	10683: 2, H0694: 2,	.0744: 2, L0747: 2,	S0031: 2, S6024: 1,	H0294: 1, H0381: 1,	H0664: 1,	H0662: 1, S0356: 1,	S0442: 1, S0358: 1,	H0411: 1,	H0586: 1,	H0036: 1,	H0309: 1,	H0121: 1,	H0081: 1, H0014: 1,	S0051: 1,	H0622: 1,
L0758: 5,	H0658: 4,	S0276: 4,	S0280: 3,	H0188: 3,	L0769: 3,	L0622: 2,	T0060: 2,	H0181: 2,	H0494: 2,	L0775: 2,	L0514: 2,	L0511: 2,	L0540: 2,	L0809: 2,	H0683: 2,	L0744: 2,	S0031: 2,	H0294: 1,	H0483: 1, H0664:	H0662: 1,	S0442: 1,	S0376: 1, H0411: 1	S6022: 1, H0586:	H0333: 1, H0036: 1	H0590: 1, H0309:	H0044: 1, H0121:	H0081: 1,	S0388: 1, S0051: 1	H0252: 1, H0622:
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L0483: 1, H0033: 1, H0111: 1, H0032: 1, H0166: 1, S0438: 1, L0762: 1, L0504: 1, L0381: 1, L0803: 1, L0375: 1, L0376: 1, L0509: 1, L0782: 1, L0544: 1, L0542: 1, L0544: 1, L0541: 1, L0543: 1, H0693: 1, S0126: 1, H0651: 1, L0777: 1, L0751: 1, L0777: 1, L0751: 1, L0777: 1, L0753: 1, H0445: 1, H0667: 1, S0194: 1 and S0398: 1.	L0774: 3, H0658: 3, L0803: 2, L0439: 2, H0619: 1, H0586:-1, H0309: 1, S0366: 1, L0659: 1 and S0052: 1.		L0748: 18, L0755: 7, H0251: 6, L0777: 6, H0658: 4, L0754: 4, L0779: 4, S0360: 3, S0276: 3, H0674: 2, L0767: 2, L0766: 2,
	Ala-37 to Ser-42, Gly-127 to Thr-134.	Arg-4 to Gly-17, Pro-39 to Lys-46, His-96 to Arg-102.	Leu-4 to Phe-9, Gly-29 to Glu-39.
	4127	4128	4130
·	3 - 503	12 - 746	1 - 705 1 - 702
	1952	1953	1954
	HPDWT55R	HPDWT56R	HPDWU47 HPDWU47R HPDWU55 HPDWU55R
	HPDWT55	HPDWT56	HPDWU47 HPDWU55

L0775: 2, L0809: 2, H0689: 2, L0757: 2, S0358: 1, S0046: 1, H0013: 1, H0427: 1, H0123: 1, H0252: 1, H0031: 1, H0617: 1, L0769: 1, L0773: 1, L0776: 1, L0773: 1, L0776: 1, L0382: 1, L0778: 1, L0382: 1, L0783: 1, L0382: 1, L0665: 1, H0144: 1, H0670: 1, S0328: 1, H0670: 1, S0328: 1, H0699: 1, L0759: 1 and	N0434: 1. L0766: 7, L0748: 6, L0771: 5, S0422: 3, H0519: 3, L0749: 3, L0756: 3, H0735: 2, H0641: 2, H0529: 2, L0659: 2, S0374: 2, L0758: 2, H0484: 1, S0442: 1, S0358: 1, S0440: 1, S0007: 1, H0632: 1, L0483: 1, H0606: 1, H0268: 1, S0150: 1, L0763: 1,
	Pro-50 to Trp-55.
	4131
	2 - 568
,	1956
·	HPDWU60R
	HPDWU60

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L0637: 1, L0764: 1, L0768: 1, L0774: 1, L0666: 1, L0809: 1, L0438: 1, H0520: 1, H0547: 1, H0658: 1, S0328: 1, H0539: 1, S0406: 1, S0027: 1, L0744: 1, L0751: 1, L0754: 1, L0593: 1, L0777: 1, L0593: 1, L0608: 1, L0593: 1,		L0740: 3, L0758: 3, S0242: 3, L0598: 2, L0766: 2, L0803: 2, H04435: 2, H0648: 2, L0748: 2, L0756: 2, L0731: 2, L0757: 2, H0170: 1, S0300: 1, H0673: 1, H0646: 1, H0674: 1, H0616: 1, L0770: 1, L0637: 1, L0794: 1, L0804: 1, L0794: 1, L0804: 1,
	Ala-51 to Tyr-56, Ala-92 to Ser-101, Ser-118 to Ile-123.	Asp-14 to Ala-19.
	4132	4133
	2 - 370	38 - 163
	1957	1958
	HPDWU63 HPDWU63R	HPFMP90R
	HPDWU63	HPFMP90

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·												•								
L0775: 1, L0784: 1, L0805: 1, L0776: 1, L0666: 1, L0664: 1, L0665: 1, L0438: 1, H0689: 1, L0439: 1, L0754: 1, L0750: 1, L0759: 1 and L0480: 1.			H0255: 2, S0114: 1, L0510: 1 and S0216: 1.			L0809: 5, L0666: 4,	L0747: 3, L0750: 3,	L0764: 2, L0803: 2,	L0774: 2, L0651: 2,	L0663: 2, L0664: 2,	L0748: 2, L0756: 2,	L0779: 2, L0755: 2,	H0669: 1, H0392: 1,	H0600: 1, S0346: 1,	H0494: 1, H0646: 1,	L0763: 1, L0371: 1,	L0372: 1, L0775: 1,	L0659: 1, L0518: 1,	L0788: 1, L0665: 1,	S0330: 1, S0392: 1,
	Phe-5 to Leu-11, Ser-20 to Arg-45.	Gly-4 to Asn-9.	Pro-11 to Lys-16, Val-18 to Arg-45, Asn-56 to Glu-67.													•				
·	4134	4135	4136	4137	4138	4139		-			,	,								
	209 - 394	3 - 182	2 - 364	2-655	114 - 212	239 - 412			•	,						- ,		,		
	1959	1960	1961	1962	1963	1964														
	HPIAL09R	HSAMY44 HSAMY44R	HSAYW59R	HSCPJ17R	HSDIA22R	HSDII65R											-			
	HPIAL09	HSAMY44	HSAYW59	HSCPJ17	HSDIA22	HSDII65												-		

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					,																						
.0749: 1, L0752: 1,	L0757: 1, S0260: 1, L0596: 1, L0591: 1 and L0608: 1.		- 1	AR033: 4, AR089: 3, AR096: 3, AR060: 3,		R055: 1, AR039: 0,	R052: 0	L0774: 3, L0752: 3,	.0592: 3, H0625: 2,	L0662: 2, L0747: 2,	,0777: 2, H0580: 1,	.0717: 1, H0455: 1,	S0280: 1, H0251: 1,	H0633: 1, L0773: 1,	.0768: 1, L0794: 1,	0803: 1, L0775: 1,	L0653: 1, L0776: 1,	L0661: 1, L0659: 1,	L0526: 1, L0809: 1,	L0665: 1, H0658: 1,	H0696: 1, L0755: 1,	L0731: 1, L0757: 1,	L0759: 1 and H0352: 1				
3	<u> </u>		Ser-20 to Ser-31.		<u> </u>	•	<u>∢</u>		.el				0	114	. <u></u>									Lys-18 to Lys-30.	+-	+	_
		4140	T																					4143	4144	11/4	414
		2 - 187	2-169	40 - 141																				40 - 180	08 6	20,	2-400
		1965	1066	1967									_											1068			1970
		+-	AC/AICAIT	┿																				010170011	HSDCJCIR	HSIFF84K	HSKJR50R
		十		HSDZE12																					HSDZJ21	HSIFF84	HSKJR50

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	H0596; 5, 80388; 4, H0598; 4, H0598; 4, S0412; 4, S0282; 3, S0280; 2, H0594; 2, H0366; 2, H0542; 2, H0395; 1, H0411; 1, H0592; 1, H0559; 1, H0427; 1, S0050; 1, H0477; 1, S0050; 1, H0436; 1, S0260; 1, S0194; 1, And H0506; 1, S0194; 1		L0755: 5, L0748: 4, H0616: 3, L0769: 3, L0752: 3, L0731: 3.	L0757: 3, S0356: 2, S0408: 2, S0046: 2,
Asn-45 to Ser-54.		Glu-77 to Arg-82, Pro-132 to Met-139.	Asp-67 to Leu-78, Asn-81 to Ser-90, Pro-96 to Glu-110, Ala-164 to Ala-171, Gln-173 to Lys-179. Ala-1 to Cys-21, Trp-23 to Ser-33. Leu-43 to Thr-58. Ser-41 to Ile-50, Gln-55 to Glu-65, Cys-69 to Asp-78.	
4146	4147	4148	4150 4151 4152 4153	
2 - 196	1 - 174	1 - 612	2 - 631 1 - 162 1 - 342 3 - 293	
1971		1973	1975 1975 1977 1978	·
HSLHT27 HSLHT27R	НЅОЪВ 16R	HSPSB24R	HSPSB74R HSPSB80R HSPSC30R HSPSC30R	
HSLHT27	HSODB16	HSPSB24	HSPSB74 HSPSB80 HSPSC30 HSPSC30	

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.0770; 2, L0771; 2,	0776: 2, L0526: 2,	0747: 2, L0750: 2,	0756: 2, L0758: 2,	,0759; 2, L0588; 2,	H0170: 1, L0459: 1,	HO484: 1, S0358: 1,	S0360: 1, H0637: 1,	S0045: 1, S0132: 1,	H0393: 1, S0278: 1,	3039: 1, H0427: 1,	10318: 1, S0049: 1,	10009: 1, H0081: 1,	10003: 1, H0051: 1,	H0083: 1, S0003: 1,	10622: 1, H0708: 1,	H0316: 1, H0038: 1,	•	S0038: 1, S0150: 1,	H0529: 1, L0762: 1,	.0763: 1, L0646: 1,	.0644: 1, L0768: 1,	0766: 1, L0774: 1,	0775: 1, L0661: 1,	0657: 1, L0782: 1,	0788: 1, L0666: 1,	,0665: 1, H0683: 1,	H0660: 1, S0378: 1,	S0406: 1, L0777: 1,	H0445: 1 and H0542: 1.
<u> </u>	<u>3</u>	<u>a</u>	2	ន	H	H	SC	<u>SC</u>	H	T	Ħ	Ä	Ĕ	Ē	Ī	臣	豆	·	重	ā	ā		7	Þ	_=	<u>.</u>	<u> </u>	Š	H
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H0124: 19, H0052: 10,	.0751: 7, L0747: 7,	H0255: 6, H0441: 6,	H0556: 5, S0222: 5,	10265: 4, S0360: 4,	10620: 4, L0655: 4,	H0144; 4, L0439; 4,	.0596: 4, L0601: 4,	S0132: 3, H0575: 3,	H0123: 3, H0012: 3,	H0024: 3, H0617: 3,	H0087: 3, H0529: 3,	.0764: 3, L0662: 3,	.0766: 3, L0663: 3,	.0565: 3, L0438: 3,	H0660: 3, L0743: 3,	.0740: 3, H0445: 3,	H0218: 2, H0585: 2,	S0418: 2, S0444: 2,	S0408: 2, H0208: 2,	S0278: 2, H0261: 2,	H0333: 2, H0706: 2,	H0581: 2, S0049: 2,	H0009: 2, L0471: 2,	S0051: 2, T0010: 2,	H0083: 2, H0594: 2,	r0006: 2, H0628: 2,	H0038: 2, S0440: 2,	S0142: 2, S0344: 2,	.5565: 2, L.5566: 2,
4154 Leu-41 to Asp-48.	,	_===						<u> </u>				. —				,	<u> </u>	<u> </u>	<u> </u>	<u> </u>		. 	<u> </u>	<u> </u>		·C		<u> </u>	I
4154										_																,			
3 - 191																													•
1979				•						,					_				_										
HSPSG03 HSPSG03R																													
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7: 2.	6: 2,	3: 2,	70: 2,	8: 2,	L0752: 2,	1: 2,	7:2,	24: 1,	+1:1,)2: 1,	[4: 1,	41: 1,	22: 1,	07: 1,	76: 1,	00: 1,	11:1,	574: 1,	22: 1,	4:1,	156: 1,)48: 1,	309: 1,	231: 1,	546: 1,	150: 1,	050: 1,	373: 1,	071: 1,
0776.2 1.0657: 2.	659: 2, L066	L0665: 2, H0683: 2,)658: 2, H067	S0146: 2, L0748: 2,	,0756: 2, L075	755: 2, L073	3758: 2, L058	L0361: 2, H0624: 1,	0002: 1, H01	0717: 1, S040	0295: 1, S01	L0785: 1, H0341: 1,	0212: 1, S028	H0722: 1, S0007: 1,	0046: 1, S047	10619: 1, S03	H0351: 1, H0411:	10392: 1, HO	10492: 1, LO6	,0623: 1, H02	H0250: 1, H0156: 1,	H0618: 1, TO	H0434: 1, H0309: 1	H0597: 1, H0231:	H0544: 1, H0546: 1,	H0046: 1, H0150: 1	H0050: 1, SO	H0016: 1, H0373:	H0051: 1, H0071:
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H0690: 1, H0682:	H0435: 1, H0539: H0521: 1, S0404: 1	\$3012: 1, \$0037: 1,	\$3014: 1, 500 \$0006: 1, 1,000	L0750: 1, L07	L0753: 1, L07	L0593: 1, S02	H0423: 1, H0506: 1,	H0008: 1 and H0293:			· •		39,	-132,	-147.		4	<u> </u>	. Š	104,	-171.				·	-156,	
										Glu-6 to Thr-11,		Thr-68 to Ile-76,	Met-80 to Arg-8	Glu-123 to Gln-132	Arg-140 to Leu-147	Glu-1 to Ile-6.	+-		Val-82 to Gln-89	Gly-99 to Ala-104,	Tyr-165 to Leu-17	 			1 4 cm 7 to 1 ve. 15		
									4155	4156						4157	4150	47.70				4159	4160	4161		4107	
	<u></u>		-						2 - 784	3 - 596						123 - 608	197 661	100-+				11 - 793	3 - 431	2 530		3 - 734	
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	H0124: 19, H0052: 10,	H0255: 6, H0441: 6,	H0556: 5, S0222: 5,	H0265: 4, S0360: 4,	H0620: 4, L0655: 4,	H0144: 4, L0439: 4,	L0596: 4, L0601: 4,	S0132: 3, H0575: 3,	H0123: 3, H0012: 3,	H0024: 3, H0617: 3,	H0087: 3, H0529: 3,	L0764: 3, L0662: 3,	L0766: 3, L0663: 3,	L0565: 3, L0438: 3,	H0660: 3, L0743: 3,	L0740: 3, H0445: 3,	H0218: 2, H0585: 2,	S0418: 2, S0444: 2,	S0408: 2, H0208: 2,	S0278: 2, H0261: 2,	H0333: 2, H0706: 2,	H0581: 2, S0049: 2,	H0009: 2, L0471: 2,	S0051: 2, T0010: 2,	H0083: 2, H0594: 2,	T0006: 2, H0628: 2,	H0038: 2, S0440: 2,	S0142: 2, S0344: 2,
Glu-181 to Phe-190.								-												-								
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1.5566: 2.		2, L0666: 2,	,0665: 2, H0683: 2	10658: 2, H0670: 2,	S0146: 2, L0748: 2,	2, 1.0752: 2		2, L0587: 2	2, H062	1, H014	1, S040	1, S011	0785: 1, H0341: 1,	1, S028	H0722: 1, S0007: 1	1, S047	1, S03(1, H0411:	1, H0574: 1	1, LOG	1, H0244:	: 1, H0156: 1	: 1, T00	1, H03		: 1, HO	: 1, HO	1, S00	H0016: 1, H0373:
5565.2	0776: 2,]	.0659: 2,	.0665: 2	H0658:	30146: 2	L0756: 2,	L0755: 2,	L0758: 2,	L0361:	T0002: 1, H0141: 1	H0717:	H0295: 1, S0114: 1,	L0785:	S0212:	H0722:	S0046:	H0619: 1, S0300: 1,	H0351:	H0392: 1,	H0492: 1	1.0623: 1	H0250: 1	H0618: 1	H0434: 1	H0597: 1.	H0544: 1	H0046: 1,	H0050: 1,	H0016
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H0520: 1, H0547: 1, H0690: 1, H0682: 1, H0435: 1, H0539: 1, H0521: 1, S0404: 1, S3012: 1, S0028: 1, S0206: 1, L0742: 1, L0750: 1, L0779: 1, L0753: 1, L0757: 1, L0593: 1, S0276: 1, H0008: 1 and H0293: 1.	H0617: 14, L0605: 14, L0657: 11, H0682: 11, H0521: 10, S0360: 8, H0620: 5, H0687: 5, L0664: 5, H0687: 5, S0406: 5, S0376: 4, H0059: 4, H0648: 4, L0768: 4, H0658: 4, H0670: 4, H0666: 4, H06713: 3, H0716: 3, H0713: 3, H0716: 3, H0618: 3, H0638: 3, H0687: 3, H0639: 3, L0662: 3, L0775: 3, L0662: 3, L0775: 3,
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H0435:	H0445:	L0581:	H0585:	T0049:	S0476:	H0550:	H0559:	H0575:	H0581:	H0606:	H0649:	L0382:	L0743:	L0599:	H0543:	H0171:	S0040:	H0650:	H0255:	H0663:	H0125:	S0442: 1,	S0358:	H0722:	S6026:	L0623:	1,0021:	T0082:	H0327:
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H0542: 1, H0422: 1, S0456: 1 and H0352: 1.			H0683: 242, S0358: 3, L0745: 3, H0656: 2, H0484: 2, S0354: 2, H0370: 2, H0427: 2, H0521: 2, S0001: 1, H0521: 2, S0001: 1, H0642: 1, H0618: 1, H0626: 1, H0628: 1, H0623: 1, H0130: 1, S0210: 1, L0438: 1, L06593: 1, H0639: 1, H06					
	Pro-6 to Gly-13, Pro-23 to Lys-32, Lys-94 to Ala-112.	Glu-17 to Asp-32.				Ala-1 to Lys-9, Val-11 to Trp-17.		Ser-1 to Thr-8,
	4165	4166	4167	4168	4169	4170	4171	4172
	1 - 339	1 - 327	2 - 442 2	2 - 406	244 - 438	2-61	46 - 210	36 - 302
	1990	1991	1992	1993	1994	5661	1996	1997
	HSPSQ22R	HSPSQ57R	HSPSY67R	HSPSZ69R		HSPTN57R	HSSDM17R	
	HSPSQ22	HSPSQ57	HSPSY67	69ZSASH	HSPTA57	HSPTN57	HSSDM17	HSWBF04

																													
			-		L0803: 10, L0794: 4,	L0779: 4, L0789: 2,	H0648: 2, S0442: 1,	S0045: 1, S0036: 1,	L0520: 1, L0763: 1,	L0769: 1, L0800: 1,	L0644: 1, L0764: 1,	L0773: 1, L0767: 1,	1 0004: 1 1 0774: 1	L0804: 1, L0774: 1,	100003: 1, 20011: 1,	1.0758: 1.	H0615: 2 and S0134: 1.								L0439: 15, H0052: 11,	S0007: 9, L0438: 6,	L0731: 6, L0779: 5,	L0754: 4, H0550: 3, 1 0769: 3, 20126: 3.	
Cl., 17 to Ale 32	410-17 to Ata-32,	Met-9 to Arg-14.	Leu-23 to Lys-29,	Lys-50 to Asn-56.	Ser-7 to Gly-12,	Pro-23 to Pro-40,	Ala-49 to Val-54.					•					Ser-4 to Thr-10,	Leu-42 to Pro-51.	4176 Am 55 to Val-63	Cys-80 to Arg-92.		Pro-8 to Trp-14,	Arg-20 to Ser-27,	Glu-52 to Glu-60.	Pro-26 to Gly-39,	Pro-68 to Arg-75.			
		1173			4174			-									4175		1196	41/0	4177	4178			4179				
			100 - 001		84 - 308							 •	•				136-207	- 001	70,	131 - 406	107 - 295	160 173	C74 - 601		94 - 366	3			
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.0743: 3, H0194: 2,	10687: 2, H0623: 2,	.0768: 2, L0776: 2,	0659: 2, L0666: 2,	0663: 2, H0689: 2,			0752: 2, L0758:	.0608: 2, H0352: 2,	H0662: 1, S0356:	S0354: 1, S0444: 1	, S04	H0441: 1, H0431:	1, HO	H0575: 1, H0590:	70048: 1, H0150:	1, S0050:	S0388: 1, H0252:	H0039: 1, H0135:	H0038: 1, H0264:			,107	3	Š		H.	, H0	H0593: 1, H0658:	H0672: 1, H0539:
743:	687: 3	768: 2	559: 2	363: 2	50330: 2, 1	0786: 2, 1	752: 2	508: 2	962:	354: 1	145:1	4	333:	575:	748: 1	H0024: 1	888: 1	039:	338:	H0494: 1	0372: 1,	0521: 1	0803: 1	0653: 1	0809: 1,	.0664: 1	.0352: 1,	593:	572:
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R0406: 1, L0751: 1	L0749: 1, L0756: 1 L0753: 1 and H050	Arg. 8 to Arg. 28	11 to A 22 10	Arg-11 to Arg-10,	Ser-42 to Arg-52,	Met-62 to Pro-73,	Pre-102 to Gly-113.				Arg-2 to Gly-12,	Alg-20 (2) 11.	y + 6 to 1 as 30;	Lys-94 to Arg-110.	Met-4 to Thr-22,	Asp-45 to Gly-50,	Gly-65 to Asp-70,	/s-96 to Asp-104,	le-121 to Leu-126,	Met-133 to Glu-144,	Ala-149 to Asn-156.	Ile-36 to Glu-41,	Glu-55 to Lys-63,	Lys-68 to Gln-81,	Pro-84 to Lys-90,	Pro-108 to Ser-119.	Pro-22 to Pro-28,	Glu-109 to Arg-118.
		4100	7	4181 Ar	Se	<u>×</u>	P. P.	4182	4183		4184 AI	₹ ₹	5 .	<u> </u>	4185 M	ď	Ö	<u>।</u>		<u>~</u>		4186 11		<u> </u>	Δ.	<u>, p.</u>	4187 P	
		+	+	19 - 390				187 - 288	351 - 620		1 - 462				94 - 618		-					208 - 717			·		2-373	
Ī		1	2002	2006			-	2007	2008		2009				2010							2011			_		2012	
			-	HUSGAIIR				HISH 88R	1UUCH08R	-	HUVFA39R		-	-	UNICA A 31R	ALCOAD VI						17 V V JUL	ALCOUNT				HVCAA68R	NA CESTA
			HULAP70	_				LITICIT 88	HUUCH08 HUUCH08R		HUVFA39	-	-		1870 4 21	HVCAA31						204 40:22	HVCAA3/				89 V V J L H L	HACAROS

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	H0666: 696, H0660:	111, L0666: 7, H0617: 4,	LU664: 4, HU349: 3,	H0620: 3, H0181: 3,	H0593: 3, H0684: 3,	H0484: 2, H0619: 2,	H0051: 2, H0606: 2,	H0670: 2, L0748: 2,	L0439: 2, S0040: 1,	H0483: 1, H0255: 1,	H0662: 1, S0007: 1,	~	H0597: 1, H0046: 1,	_:	`	H0517: 1, L0770: 1,	•	L0783: 1, L0809: 1,	H0435: 1, H0658: 1,	H0539: 1, H0521: 1,	H0696: 1 and S0434: 1.						L0769: 7, L0747: 7,	L0777: 7, L0752: 7,	L0665: 6, L0770: 5,
						-								•			•		:			Arg-1 to Gln-18,	Asp-52 to Leu-57,	Asp-67 to Lys-74,	Asp-85 to Ala-92.				,
4188	4189											•										4190				4191	4192		
1 - 462	2 - 649																		•			353 - 664				3 - 758	3 - 224		
2013	2014																					2015				2016	2017		
HVCAA94R	HVCAB02R																•					HVCAB03R				HVCAB18R	HVCAB44R		•
HVCAA94	HVCAB02																					HVCAB03				HVCAB18			

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L0766: 5, L0771: 4, L0666: 4, L0771: 4, L0666: 4, L0771: 4, L0666: 4, L0771: 3, L0762: 3, L0763: 3, L0743: 3, L0749: 3, L0761: 2, L0769: 1, R0661: 1, H0402: 1, R0661: 1, H0409: 1, R0661: 1, H0699: 1, L0777: 1, H0409: 1, L0777: 1, H0409: 1, L0777: 1, H0409: 1, H0678: 1, H0679:		
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			Lys-6 to Thr-11, Gly-22 to Phe-30, Leu-40 to Gly-53,	Arg-56 to Lys-64, Lys-93 to Leu-101, Met-176 to Leu-181	Arg-7 to Asp-13.	
	4193	4194	4195		4196	4197
	4 - 612	1 - 357	83 - 739		319 - 417	1 - 369
	2018	2019	2020		2021	
	HVCAB52R	HVCAB57R	HVCAB88R		HVCAC42R	HVCAD52 HVCAD52R
	HVCAB52	HVCAB57	HVCAB88		HVCAC42	HVCAD52

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Tyr-44 to Asp-49, Pro-132 to Leu-139, Cys-168 to His-174.	Pro-49 to Val-58, Gln-60 to Gln-78, Arg-85 to His-99, Phe-122 to Asp-128, Arg-157 to Arg-163,	O E D & D & D E E & D H H H H F & & & & & & & & & & & & & & &
4198 Tyr-4 Pro-13 Cys-1	4199 Pro-4 Gln-6 Arg-8 Phe-1 Arg-1	4200 Gly-1
49 - 663	3 - 605	2 - 493
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	Asn-20 to Gln-27.	Arg-92 to Arg-101, Lys-134 to Asn-140 Ser-149 to Thr-159 Gly-166 to Ile-175.
	Asn-2	Arg-9 Lys-1 Ser-14 Gly-1
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H0617:	H0166:	H0163:	H0038:	L0564:	T0042:	S0438:	H0641	L0762: 1	L0372: 1,]	1,0771	L0521: 1, 1	99201	1 5574: 1	1 0650	I 0276. 1	7.050 T		1.0518: 1	L0545	L0666: 1	50216	H059	690H	990H	H053	8033	H070	8014	1074	H044
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L0596: 1, L0588: 1, L0599: 1, L0366: 1, S0011: 1, H0668: 1, H0216: 1, S0242: 1, S0196: 1, H0542: 1, H0543: 1, L0697: 1, L0698: 1, S0398: 1, S0424: 1, S3020: 1 and S0460: 1.		-	S0408: 5, L0809: 5,	L0800: 4, L0804: 4,	H0647: 3, L0803: 3,	L0775: 3, L0805: 3,	S0126: 3, L0751: 3,	S0360: 2, S0278: 2,	H0409: 2, H0424: 2,	H0617: 2, H0494: 2,	L0783: 2, H0670: 2,	L0747: 2, L0777: 2,	L0731: 2, L0601: 2,	H0556: 1, H0295: 1,	S0116: 1, H0341: 1,	S0442: 1, S0358: 1,	S0376: 1, H0370: 1,	H0042: 1, H0575: 1,	S0049: 1, H0150: 1,	H0163: 1, S0294: 1,	S0438: 1, H0633: 1,
	Asn-1 to Lys-6.				-																
	4208	4209	4210																•		
·	108 - 365	2 - 508	1 - 537		-																
	2033	2034	2035				•														
	HVCAI24 HVCAI24R	HVCAI79R	HVCAJ27R										٠		<u>,</u>			-			
	HVCA124	HVCAI79	HVCAJ27												·						

H0529: 1, L0769: 1, L0374: 1, L0764: 1, L0560: 1, L0774: 1, L0376: 1, L0629: 1, L0667: 1, L0657: 1, L0666: 1, H0682: 1, H0683: 1, H0658: 1, H0666: 1, S0328: 1, H0539: 1, H0555: 1, L0779: 1, L0752: 1, L0779: 1, L0752: 1, L0757: 1, L0758: 1, L0759: 1, H0667: 1 and H0423: 1.				T10000. 2 T10125. 3	H0009: 3, H0153: 3, H0494: 3, H0265: 2, H0341: 2, H0484: 2,	S0046: 2, S0222: 2,	H0266: 2, 50144: 2, H0658: 2, L0742: 2,	H0542: 2, H0423: 2,	1101/4:4:4:4:4:4:
	Lys-61 to Gln-73, Gln-81 to Ser-97.	Arg-14 to Gly-41, Arg-47 to Glu-53, Gln-56 to Lys-70.	Glu-1 to Lys-9.						
	4211	4212	4213	4214	4215				
	1-477	1 - 210	3 - 467	3 - 383	1-516				
	2036	2037	2038	2039	2040				
	HVCAJ81R	HVCAJ95R	HVCAK02R	HVCAL06R	HVCAO17 HVCAO17R		<u>.</u>		
	HVCAJ81	HVCAJ95	TYCAKO	HVCAI 06	HVCA017				

H0685: 1, S0040: 1, H0295: 1, S0420: 1, S0356: 1, S0420: 1, H0392: 1, H0586: 1, H0599: 1, H0618: 1, H0544: 1, H0612: 1, H0024: 1, H0012: 1, H0024: 1, H0012: 1, H0053: 1, H0040: 1, H0063: 1, H0640: 1, H0060: 1, H0640: 1, H0560: 1, H0670: 1, H0560: 1, H0670: 1, H0668: 1, L0599: 1, L0757: 1, L0599: 1,	S0026: 1, H0667: 1, H0136: 1, S0276: 1 and H0543: 1.			
		Pro-56 to Thr-65.		Val-105 to Ile-113,
	, .	4216	4217	4218
		10 - 306	3 - 407	28 - 480
		2041	2042	2043
		HVCA063R	HVCAP89R	HVCAQ53R
		HVCA063		HVCAQ53 HVCAQ53

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				L0777: 12, H0341: 6,	S0440: 6, S0126: 6,	S0358: 5, H0250: 5,	H0428: 5, H0031: 5,	H0090: 5, L0747: 5,	.0731: 5, H0542: 5,	S0422: 4, H0547: 4,	t8: 4, L0749: 4,	.0759: 4, H0624: 3,	H0580: 3, H0486: 3,	S0003: 3, S0214: 3,	H0494: 3, L0770: 3,	.0764: 3, L0758: 3,	99: 3, S0026: 3,	H0543: 3, H0170: 2,	H0657: 2, H0656: 2,	S0420: 2, S0442: 2,	S0444: 2, S0010: 2,	H0318: 2, H0581: 2,	H0046: 2, H0014: 2,	H0083; 2, L0483; 2,	H0040: 2, H0551: 2,	0769: 2, L0773: 2,	.0766: 2, L0774: 2,	.0659: 2, H0144: 2,
Asp-128 to Phe-137, Arg-145 to Arg-151.	Asp-15 to Thr-23.	-	·		8044	8035	H042	500H	1,073	S042	7.07	107	H05	008	H04	100	105	HOS	H06	804	S04	HO	HOC	OH	OH			TO
	-1		4221	4222					_																		.	
	193 - 402	3 - 455	2-715	1	•																							
	2044	2045	2046		1																							
	HVCA084R	HVCAR87R	HVCASORR	HAVOA SKOB	HVCASSA			•																		_		
	HVCAO84	HVCAR87 HVCAR87R	111/0/00/00	HVCASUS	HVCAS52																							

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L0565: 2, H0659: 2.	H0666: 2, H0648: 2,	S0328: 2, H0525: 2,	10696: 2, H0134: 2,	30406: 2, L0756: 2,	2, L0757: 2,	S0436: 2, L0595: 2,	S0384: 2, H0506: 2,	H0171: 1, S0040: 1,	.0049: 1, H0661: 1,	H0638: 1, S0418: 1,	S0354: 1, S0360: 1,	S0410: 1, S0007: 1,	S0300: 1, S0278: 1,	S6014: 1, H0431: 1,	H0587: 1, H0333: 1,	HO574: 1, HO270: 1,	.0040: 1, L0586: 1,	H0635: 1, H0156: 1,	70082: 1, H0706: 1,	S0346: 1, H0421: 1,	H0052: 1, H0597: 1,	H0545: 1, H0242: 1,	.0471: 1, H0373: 1,	S0051: 1, H0375: 1,	H0594: 1, H0687: 1,	HO288: 1, HO328: 1,	H0553: 1, L0143: 1,	H0032: 1, H0598: 1,	H0634: 1, H0616: 1,
1.0565	9990H	S0328:	9690H	S0406	L0752	S0436	S0384:	H0171	T0049	H0638	S0354:	S0410:	S0300	S6014:	H0587	H0574	T0040	H0635	T0082	S0346:	H0052	H0545	L0471:	S0051:	H0594	H0288	H0553	H0032	H0634
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H0413: 1, H0100: 1,	T0042: 1, H0560: 1,	S0150: 1, H0633: 1,	H0652: 1, L0763: 1,	L0371: 1, L0638: 1,	L0667: 1, L0648: 1,	L0521: 1, L0662: 1,	L0649: 1, L0803: 1,	L0776: 1, L0783: 1,	L0383: 1, L0647: 1,	L0664: 1, L0665: 1,	H0520: 1, H0689: 1,	H0414: 1, H0658: 1,	H0670: 1, H0660: 1,	H0710: 1, S0152: 1,	S0332: 1, S0454: 1,	H0522: 1, H0704: 1,	H0214: 1, S3012: 1,	S3014: 1, S0027: 1,	S0028: 1, S0206: 1,	L0740: 1, L0750: 1,	S0434: 1, L0581: 1,	H0653: 1, H0667: 1,	S0242: 1, S0194: 1,	H0422: 1 and S0424: 1.	L0748: 6, L0439: 6,	L0731: 6, H0663: 4,	S0354: 4, H0038: 3,	S0150: 3, L0666: 3,	
				 -																					7007	4772			
						_																				114 - 257			
-								-																		2048			
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58: 3,	197: 2,	112: 2,	10: 2,	26: 2,	561: 1,	20: 1,	08: 1,	131: 1,	40: 1,	13: 1,	90: 1,	52: 1,	73: 1,	12: 1,	22: 1,	53: 1,	113: 1,	40: 1,	7. 1.	89: 1,	59: 1,	89: 1,	97: 1,	20: 1,	59: 1,	52: 1,	36: 1,	12: 1,	5: 1,
S0404: 3, L0758: 3,	S0212: 2, H0497: 2,	H0551: 2, H0412: 2	H0561: 2, S0210: 2,	.0521: 2, L0526: 2,	H0506: 2, H0661: 1	S0418: 1, S0420: 1	: 1, SO4	10619: 1, H0431:	H0486: 1, T0040:	70060: 1, H0013:	.0021: 1, H0590:	10581: 1, H0052:	H0263: 1, H0373:	H0266: 1, S0312:	S0314: 1, S0022:	H0622: 1, H0553: 1	H0708: 1, H0413: 1	H0560: 1, L0640: 1	1, L0764:	1, L0389:	1, L0659:	1, L0789:	1, H0697:	S0374: 1, H0520:	H0435: 1, H0659:	H0666: 1, S0152:	S0406: 1, H0436: 1	1, L0742:	.0753: 1, L0595:
S0404	S0212	H0551	H0561	L0521	H0506	S0418	S0358	H0619	H0486	T0060	17007	H0581	H0263	H0266	S0314	H0622	H0708	H0560	L0646: 1,	L0649: 1,	L0804: 1,	L0543: 1,	L0664: 1,	S0374:	H0435	9990Н	S0406 :	S0027: 1,]	L0753:
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H0668: 1, S0276: 1 and H0422: 1.		-			410 4 0006.	ARU39: 419, ARU30. 339, AR053: 225,	AR052: 212, AR104: 1168 AR089: 168.	AR060: 114, AR033:	108, AR055: 99, AR061: 74												
	Arg-7 to Val-15.	Val-3 to Lys-9, Leu-32 to Gln-40.								•	Lys-7 to Thr-13, Asp-24 to Thr-30,	Gly-39 to Glu-52.					Ser-25 to Asn-31,	Asp-38 to Val-48, Ser-57 to Glv-63.		Pro-1 to Ser-7,	Val-22 to Asn-29,
	4224	4225	4226	4227	4228	4229					4230		4231	4232	4233	4234	4235		4236	4237	
	3 - 404	190 - 309	3 - 431	3 - 524	3 - 251	2 - 361					22 - 318		2 - 448	2 - 298	34 - 699	1 - 156	203 - 469		1 - 315	31 - 459	
	2049		2051	2052	2053	2054					2055		2056	2057	2058	2059	2060		1906	2062	
	HVCAU64R	 	HVCRE76R	+-	HVCBF38R	HVCBF89R					HVCBG01R		HVCB031R	HVCCA08R	HVCCK34R	HVCCR91R	HVCCV93R		dar A dorar	HVCDA/5 HVCDA/3R	
	HVCATI64	HVCBD18	7/CBE7/VI	_	HVCBF38	HVCBF89					HVCBG01		HVCBO31	1	HVCCK34	HVCCR91	HVCCV93			HVCDA/5	מעטאה

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			H0666: 8, S0144: 6,	L0747: 5, H0545: 4,	L0769: 4, L0766: 4,	L0731: 4, H0341: 3,	L0764: 3, L0794: 3,		L0758: 3, S0045: 2,	H0257: 2, H0581: 2,	H0546: 2, H0135: 2,	L0803: 2, L0657: 2,	L0789: 2, L0666: 2,	S0044: 2, H0445: 2,	L0588: 2, L0594: 2,	L0601:2, H0543:2,	S0040: 1, H0295: 1,	H0583: 1, S0116: 1,	S0212: 1, H0661: 1,	H0664: 1, S0418: 1,	S0356: 1, S0358: 1,	S0360: 1, S0007: 1,	S0046: 1, S0132: 1,	S0278: 1, H0592: 1,	H0586: 1, H0559: 1,	H0575: 1, S0182: 1,	H0620: 1, S0388: 1,	H0083: 1, S0312: 1,
Pro-45 to Pro-51,	Arg-59 to Val-64,	1 di / t to ta / 2 / 2 / 2 / 2	Ser-56 to Phe-69,	Arg-113 to Thr-123,	Leu-128 to Ser-135,	Lys-148 to Ala-156,	Gly-166 to Ser-171,	Glu-188 to Leu-194,	Ala-211 to Gly-222.	•					•													
		4238	4239																									
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		2063																	_									
		HVCDF50R	HVCDH77 HVCDH77R																									
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H0617: 1, H0169: 1, H0708: 1, H0163: 1, H0551: 1, H0488: 1, T0069: 1, H0102: 1, H0100: 1, H0494: 1, L0475: 1, H0646: 1, S0142: 1, S0002: 1, L0762: 1, L0796: 1, L0646: 1, L0796: 1, L065: 1, L0771: 1, L0783: 1, L0790: 1, L0783: 1, L0790: 1, H0144: 1, L0438: 1, H0519: 1, H0134: 1, S3014: 1, L0790: 1, L0779: 1, L0790: 1, L0779: 1, L0790: 1, S0031: 1, L0592: 1, H0665: 1, S0276: 1 and H0665: 1, S0276: 1 and				L0766: 12, S0412: 8, S0422: 4, L0756: 4, H0341: 3, H0638: 3, L0770: 3, L0666: 3, L0777: 3, S0360: 2,	HU316: 2, HU32/: 2,
	3 - 155 4240	74 - 619 4241	56 - 238 4242	403 - 570 4243	_
·	/VAB25R 2065	/VAB37R 2066	VVAC18R 2067	/VAD74R 2068	
	HVVAB25 HV	HVVAB37 HV	HVVAC18 HV	HVVAD74 HVVAD74R 2	

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.0803: 2, H0670: 2,	,0740: 2, L0731: 2,	S0260: 2, S0436: 2,	H0170: 1, H0125: 1,	S0418: 1, S0444: 1,	10489: 1, H0580: 1,	H0734: 1, S0222: 1,	H0587: 1, H0497: 1,	H0013: 1, H0075: 1,	H0427: 1, H0046: 1,	H0354: 1, L0142: 1,	.0055: 1, H0090: 1,		H0647: 1, H0529: 1,	.0369: 1, L0371: 1,	.0772: 1, L0646: 1,		.0381: 1, L0650: 1,	L0774: 1, L0653: 1,	.0655: 1, L0606: 1,		.0663: 1, L.0664: 1,	S0374: 1, H0593: 1,	H0711: 1, H0659: 1,	H0658: 1, H0672: 1,	H0521: 1, L0779: 1,	_		S0242: 1, H0423: 1 and	H0721: 1.
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	H0672: 2, L0748: 2, H0156: 1 and H0494: 1.	L0747: 5, L0731: 4, L0742: 3, L0745: 3, L0766: 2, L0766: 2, L0766: 2, L0604: 2, T0049: 1, S0258: 1, H0549: 1, H0263: 1, H0551: 1, H0563: 1, H0571: 1, L0771: 1, L0768: 1, L066: 1, L0677: 1, L066: 1, L0677: 1, H0648: 1, H0672: 1, H0648: 1, H0672: 1, H0648: 1, L0779: 1, L0769: 1, L0779: 1, L0750: 1, L0779: 1, L0750: 1, L0779: 1, L0750: 1, L0779: 1, L07780: 1 and L0592: 1.					
Glu-21 to Ile-26.	Ser-1 to Asp-9, Lys-26 to Gln-36.			Gin-33 to Ala-38, Thr-47 to Asn-64,	Pro-117 to Leu-122, Thr-160 to Gly-171.	His-1 to Trp-6, Gly-28 to Asp-34,	Arg-41 to Ala-53,
4244	4245	4246	4247	4248		4249	
96 - 347	157 - 381	3 - 245	2 - 307	3 - 527		1 - 327	·
2069	 	2071	2072	2073		2074	
	HVVAE88R	HVVAH91R		HVVAJ23R		HVVAJ28R	
UNIVA E73	HVVAE88	нууланы	HVVA103	HVVAJ23		HVVAJ28	

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	L0659: 55, H0644: 23,	L0747: 17, H0622: 16,	S0194: 16, H0553: 15,	L0662: 15, S0126: 15,	L0740: 15, L0754: 14,	H0031: 13, H0593: 12,	L0649: 11, H0435: 11,	L0757: 11, H0124: 10,	S0250: 9, S0356: 8,	H0672: 7, L0362: 7,	L0779: 6, S0210: 5,	L0771: 5, L0663: 5,	L0664: 5, H0547: 5,	L0750: 5, L0601: 5,	S0360: 4, H0619: 4,	L0764: 4, L0804: 4,	L0775: 4, S0358: 3,	H0486: 3, H0628: 3,	H0264: 3, L0451: 3,	L0776: 3, L0655: 3,	L0783: 3, H0693: 3,	H0660: 3, S0454: 3,	L0749: 3, L0605: 3,	S0192: 3, H0170: 2,	H0294: 2, S0180: 2,	H0586: 2, H0013: 2,	H0687: 2, H0428: 2,
Arg-69 to Thr-76, Leu-97 to Glu-103.			•				•					-							-								
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0483; 2, H0488; 2,	S0450: 2, H0647: 2,	0208: 2, L0648: 2,	L0523: 2, L0809: 2,	0665: 2, H0144: 2,	.0565: 2, H0690: 2,	H0518: 2, H0696: 2,	3012: 2, L0752: 2,	L0755: 2, L0731: 2,	0665: 2, H0171: 1,	H0656: 1, S0212: 1,	0376: 1, S0410: 1,	S0132: 1, H0393: 1,	0717: 1, H0392: 1,	H0331: 1, H0485: 1,	.0040: 1, T0060: 1,	0021: 1, H0599: 1,	T0082: 1, H0618: 1,	H0052: 1, H0085: 1,	70110: 1, H0046: 1,	10050: 1, L0471: 1,	40355: 1, S0003: 1,	H0615: 1, H0039: 1,	0143: 1, H0111: 1,	0455: 1, S0366: 1,	H0135: 1, H0040: 1,	H0551: 1, H0100: 1,	L0564: 1, H0509: 1,	H0517: 1, L0763: 1,	L0769: 1, L0772: 1,
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L0646: 1, L0768: 1, L0375: 1, L0784: 1, L0806: 1, L0805: 1, L0653: 1, L0654: 1, L0517: 1, L0384: 1, L0789: 1, L0791: 1, L0438: 1, H0520: 1, H0689: 1, H0682: 1, H0689: 1, H0520: 1, H0704: 1, S0152: 1, S0432: 1, S014: 1, S0027: 1, S0028: 1, L0744: 1, L0758: 1, H0595: 1, H0707: 1, S0436: 1, L0591: 1, L0599: 1, S0196: 1, H0422: 1, S0384: 1 and			L0747: 22, L0599: 11, L0766: 9, L0748: 9, L0483: 8, L0740: 7, L0665: 5, L0754: 5,
		Lys-1 to Lys-14, Phe-21 to Trp-26, Val-87 to Cys-92, Thr-139 to Gln-153, Gln-156 to Lys-162, Glu-171 to Leu-177.	·
	4251	4252	4253
·	121 - 312	3 - 782	372 - 584
	2076	2077	2078
	HVVAK42R	HVVAK46 HVVAK46R	HVVAK68R
	HVVAK42	HVVAK46	HVVAK68

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.0764: 4, L0775: 4,	,0362: 4, H0695: 3,)663: 3, H0144: 3,	,0755: 3, H0170: 2,	J0717: 2, H0437: 2,	10036: 2, H0251: 2,	10545: 2, L0471: 2,	10266: 2, H0553: 2,	H0413: 2, L0763: 2,	.0772: 2, L0655: 2,	0543: 2, L0666: 2,	10547: 2, S0406: 2,	,0745: 2, L0749: 2,	.0731: 2, L0757: 2,	.0758: 2, H0445: 2,	.0596: 2, L0591: 2,	.0581: 2, L0608: 2,	S0192: 2, H0624: 1,	S0134: 1, H0650: 1,	S0116: 1, H0663: 1,	S0356: 1, S0444: 1,	H0580: 1, S0300: 1,	H0441: 1, H0442: 1,	H0586: 1, H0587: 1,	H0333: 1, H0643: 1,	H0632: 1, H0486: 1,	H0013: 1, H0118: 1,	H0042: 1, H0004: 1,	r0048: 1, H0318: 1,	H0421: 1, H0041: 1,
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H0014: 1, S0051: 1,	H0687: 1, S0003: 1,	S0214: 1, H0688: 1,	H0031: 1, H0163: 1,	H0038: 1, H0040: 1,	 H0059: 1, T0069: 1,	L0475: 1, S0015: 1,	H0633: 1, H0538: 1,	L0769: 1, L0796: 1,	L0637: 1, L0373: 1,	L0521: 1, L0375: 1,	 L0527: 1, L0656: 1,	•	-	H0682: 1, H0435: 1,	H0659: 1, H0670: 1,	H0672: 1, S0330: 1,	H0521: 1, H0696: 1,	H0134: 1, H0214: 1,	H0555: 1, L0746: 1,	L0750: 1, L0752: 1,	L0759: 1, H0343: 1,	L0589: 1, L0590: 1,	S0194: 1, H0543: 1,	S0452: 1 and H0506: 1.		AR089: 67, AR104: 8,
	 -									-									-							Gly-72 to Asp-81.
																									4254	4255
														-			-		·						2 - 373	97 - 408
			•																	·					2079	2080
																		•							HVVAK85 HVVAK85R	HVVAL81R
																						•••			HVVAK85	HVVAL81

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AR096: 5, AR060: 4, AR055: 4, AR061: 3 H0169: 21, H0674: 19, L0754: 19, H0309: 15,	F0060: 7, H0038: 7, H0509: 7, S0044: 7, F0358: 6, H0124: 6	10163; 6, H0672; 6, 10731; 6, L0750; 5,	0603: 5, H0032: 4,	10616: 4, S0330: 4, 20749: 4, L0755: 4,	H0597: 3, S0318: 3,	H0622: 3, L0540: 3, S0328: 3, S0406: 3,	.0758: 3, S0444: 2,	10486: 2, H0122: 2,	H0355: 2, S0316: 2,	_0770: 2, L0646: 2,	.0771: 2, L0662: 2,	0364: 2, L0803: 2,	.0783: 2, H0696: 2,	.0/43: 2, LU/4/: 2,	00581: 2, F10506: 2,	H0675: 1. H0643: 1.	H0706: 1, H0253: 1,	H0085: 1, H0204: 1,	H0510; 1, S0314: 1,	H0553: 1, H0673: 1,
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S0366: 1, H0380: 1, T0004: 1, S0438: 1, L0372: 1, L0773: 1, L0542: 1, L0806: 1, L0519: 1, L0529: 1, H0691: 1, S0148: 1, H0660: 1, S0004: 1, S0146: 1, L0780: 1,						H0672: 2, S0007: 1, -0717: 1, H0545: 1,	.0764: 1, L0649: 1,	.0659: 1, L0809: 1,	.0438: 1, H0555: 1, .0748: 1, L0754: 1 and	.0752: 1.	L0659: 14, L0740: 11,	L0666: 8, H0059: 7,	H0265: 6, H0040: 6,
<u>NEUTHURSS</u>	Arg-1 to Cys-10, Asn-29 to Phe-38.		Lys-62 to Tyr-73.		Leu-36 to Ser-43, Gly-68 to Phe-73.	Lys-17 to Tyr-22.	H	<u> -</u> -	<u>-11</u> :			i ⊢ i	Li.
	4256	4257	4258	4259	4260	4261				ì	4262		
	124 - 354	3 - 413	1 - 324	2 - 436	3 - 236	7 - 171					1 - 189		
	2081	2082	2083	2084	2085	2086			٠,		2087		
	HVVAP65 HVVAP65RP	HVVAS27 HVVAS27RP 2082 00B	HVVAW26R	HVVBD91RP 00B	HVVBF09R	HVVBF66R					HVVBH88R		•
	S94VVAH	HVVAS27	HVVAW26	HVVBD91	HVVBF09	HVVBF66					HVVBH88		

3356: 5, 0539: 5, 0750: 5	0759: 4, 10575: 3,	0010: 3, 10031: 3, 10433: 3	10509: 3,	.03/2: 3, .0806: 3,	L0663: 3,	20665: 3,	JOS91: 3, 1	10587: 2,	H0036: 2,	S0049: 2,	T0110: 2,	H0123: 2,	H0024: 2, H0083: 2,	H0179: 2, T0006: 2,	50000: 2, IXO338: 2, 50000: 2, 1,0769: 2,	0796; 2, L0646; 2,	0374: 2, L0648: 2,	0776: 2, H0520: 2,	H0519: 2, H0689: 2,	10555: 2, 50028: 2,	.0439: 2, L0752: 2,
1.0596: 6, S0356: 5, S0356: 5, S0358: 5, H0539: 5, 1 0750: 5	H0431: 3, L0759: 3, H0431: 4, L0759: 4, H0442: 3, H0575: 3,	H0428: 3, H0031: 3, H0031: 3, H0031: 3, H0031: 3	H0135: 3, H0435: 3, H0412: 3, H0509: 3,	L0637: 3, L0372: 3, L0521: 3, L0806: 3,	L0519: 3, I	L0664: 3, L0665: 3,	\$0378: 3, LU591: 3,	503401 5, 110465; 2, 1104655; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2, 110465; 2,	H0013: 2, H0036: 2,	S0010: 2, S0049: 2,	H0194: 2, T0110: 2,	H0046: 2, H0123:	H0024: 2,	H0179: 2,	H0038: 2,	r 0796: 2,	1.0374: 2,	L0776: 2,	H0519: 2	H0555: 2	L0439: 2,
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L0366:	H0556	S0114;	S0212:	H0125	S0354:	H0675:	S0468:	S0005:	T0109	H0505:	H0581	H0309	H0545	L0471:	H0620:	S0051:	H0266:	S0214: 1	H0039:	H0424	H0553	H0316	, H0634:	H0551	T0041:	L0475:	S0144:	S0344:	L0638:
L0755: 2, L0366: 2	H0542: 2, H0556: 1	S0040: 1, S0114: 1	70049: 1, S0212: 1	H0638: 1, H0125:	S0420: 1, S0354:	S0376: 1, H0675: 1	H0489: 1, S0468:	S0132: 1, S0005:	H0492: 1, T0109:	S0346: 1, H0505:	H0318: 1, H0581:	H0196: 1, H0309:	H0263: 1, H0545:	10050: 1, L0471:	H0012: 1,	H0014: 1, S0051:	H0510: 1, I	H0188: 1,	H0252: 1,	H0622: 1, H0424:	H0030: 1, H0553:	H0644: 1, H0316:	50036: 1,	H0087: 1, H0551:	.0564: 1, T0041:	T0042: 1, L0475:	H0130: 1, S0144: 1	S0142: 1, S0344: 1	S0210: 1, L0638: 1
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L0761: 1, L0803: 1, L0653: 1, L0661: 1, L0527: 1, L0636: 1, L0783: 1, L0809: 1, T0068: 1, L0438: 1, S0126: 1, H0690: 1, H0682: 1, H0670: 1, H0658: 1, H0670: 1, H0651: 1, S0380: 1, H0651: 1, S0380: 1, H0651: 1, L0742: 1, L0748: 1, L0747: 1, L0756: 1, L0747: 1, L0758: 1, L0747: 1, L0758: 1, L0603: 1, S0011: 1, S0194: 1, S0196: 1, S0194: 1, S0196: 1, and H0506: 1.		·		
		Gln-5 to Pro-13, Pro-25 to Lys-32, Asp-62 to Ser-72, Gln-78 to Ser-85, Pro-91 to Glu-105.		
	4263	4264	4265	4266
	2 - 592	150 - 578	3 - 413	1 - 633
	2088	2089	2090	2091
	HVVBI16R	1	HVVBJ55 HVVBJ55RP	HV
	HVVRI16	HVVBJ08	HVVBJ55	HVVBK13

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4267 Pro-62 to Pro-73, Thr-80 to Ser-92,	Gln-108 to Tyr-116.		·	Asn-15 to Leu-27,	Lys-40 to Yat-33.		Pro-1 to Phe-9,	Ser-111 to Asn-117,	Ser-159 to Pro-167.	Ser-56 to Tyr-62,	Gln-67 to Pro-74,	Thr-83 to Gly-94,	Gln-120 to Gln-130,	Pro-150 to Ser-157.		Lys-40 to Gly-47.								****	
4267	•	4268	4269	4270	4271	4272	4273			4274					4275	4276	4277	4278							
3 - 644		50 - 433	2 - 406	116 - 316	454 - 2	3 - 161	2 - 580			1 - 792					299 - 427	224 - 481	2-640	447 - 695							
2092		2093	2094	2095	2096					2099		٠			2100	2101	2102	2103							
HVVBO65 HVVBO65R 2092		HVVBO88R		HVVBT33R	HVVRT60R					HVVCB08R					HVVCC06R	2	HVVCD81R	HVVCD87R							•
HVVBO65		HVVBO88	HVVBR70	HVVBT33	HVVRT60	HVVBY87	HVVCB04			HVVCB08					HVVCC06	HVVCD65	HVVCD81 HVVCD81	HVVCD87							

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749:	1222:	0435:	758:	3013:	0150:	0038:	662:)663:	0672:	3404:	0265:	3650:	0360:	0333:	0318	0620	0375:	0039	0040	0412	10529	0775:	0547	0436	0752:	0624	10583	0663	10638
6, L	6, 80	5, H	5, LC	5, H(4, H	4, H	4, T	4, Ľ	4, H	4, S(4, H	3, H	. 3, S.	3, H	: 3, H	: 3, H	3, H	:3, H	: 3, H	: 3, H	: 3, H	3, L	3, H	3, H	3, L	3, H	: 2, E	: 2, H	2, F
H0539: 6, L0749: 6,	.0596: 6, S0222: 5,	H0457: 5, H0435: 5,	0756:	,0601: 5, H0013: 4,	0046	H0266: 4, H0038: 4,	.0771: 4, L0662: 4,	0659:	0670	H0521: 4, S0404: 4,	J0745: 4, H0265: 3,	S0114: 3, H0650: 3,	10662	S0045: 3, H0333: 3,	H0156: 3, H0318: 3	0000	T0010: 3, H0375: 3,	H0428: 3, H0039: 3,	H0591: 3, H0040: 3,	10063	H0100: 3, H0529: 3,	.0770: 3, L0775: 3,	980	S0126: 3, H0436: 3,	J0741: 3, L0752: 3,	S0026: 3, H0624: 2,	H0713: 2, H0583: 2,	S0116: 2, H0663: 2,	H0305: 2, H0638: 2,
H	<u> </u>	프	<u> </u>	<u>ب</u>	正	프	ᄀ	<u> </u>	五	巫	ᆚ	S	<u> </u>	S	<u>正</u>	Œ	H	<u>;11;</u>	<u>;11</u> ;	بلز	<u>ili</u>	<u></u>	<u></u>	<u>(7)</u>	<u></u>	<u>.</u>	<u></u>	<u>(7)</u>	
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S0418: 2, S0442: 2,	H0580: 2, S0007: 2,	2, L0717:	10261: 2, H0392:	10586: 2, H0587:	2, H0271:	10687: 2, H0553: 2,	.0456: 2, H0616: 2,	HO413: 2, H0560: 2,	.0764: 2, L0773: 2,	, L0774:	L0378: 2, L0806: 2,	, L0517:	2, H0690	10659: 2, H0658: 2,	2, L0757:	2, H0542	2, L0600	1, S0218	H0657: 1, H0656: 1,	H0341: 1, S0298: 1,	S0212: 1, S0110: 1,	1, \$0030:	H0661: 1, S0420: 1	S0356: 1, S0376:	S0444: 1, S0046: 1	30476: 1, H0550: 1	H0441: 1, H0438: 1,	H0492: 1, H0485: 1,	H0427: 1, S0280: 1,
S0418: 2	H0580: 2	H0619: 2	H0261: 2	H0586: 2	H0635: 2,	H0687: 2	1.0456: 2	H0413: 2	L0764: 2	L0521: 2	L0378: 2	L0493: 2	10647:2	H0659: 2	L0740: 2	L0588: 2	H0422: 2, L0600: 2,	H0171:	H0657:	H0341:	S0212: 1	S0001: 1	H0661:	S0356: 1	S0444: 1	S0476: 1	H0441:	H0492:	H0427:
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30665:	1, S0049: 1,	H0251	H0263	1, H0041: 1	H0081	H0023: 1	S0051	H0267	S0250	H0252: 1, H0615: 1,	70023: 1, T0006: 1	H0031: 1, H0644:	H0673	S0364: 1, L0455: 1	H0400	S0036: 1, H0087:	10551: 1, H0102:	r0041: 1, H0625:	S0344	I, S0426: 1	L0769	L0772: 1	L0372	1.0767:	L0560	1.0375	10807	L0519: 1	S0428:
S0010: 1, S0665: 1	81: 1,	H0052: 1,	09: 1,	H0597: 1,	H0563: 1,	H0012: 1,	51: 1,	56: 1,	79: 1,	52: 1,	23: 1,	31: 1,	55: 1,	64: 1,	68: 1,	36: 1,	51: 1,	41: 1,	41:1,	S0002: 1,	.0369: 1,]	.5575: 1,	73: 1,	•	்ட்	,0650: 1,	,0661: 1,		.0792: 1,
5001	H0581: 1	HOO	H03	HOS	H05	H00	H00	H03	H01	H02	T00	H00	8	<u>803</u>	H00	<u>200</u>	H05	<u>6</u>	HO	<u>800</u>	<u> </u>	1.55	<u> </u>	<u> </u>	<u>2</u>	<u>1</u>	<u>3</u>	<u> </u>	5
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H0144: 1, L0352: 1, H0520: 1, H0682: 1, H0660: 1, H0651: 1, H0710: 1, H0518: 1, S0152: 1, H0579: 1, H0696: 1, S0146: 1, H0676: 1, R0031: 1, S3012: 1, S0037: 1, L077: 1, L0779: 1, L077: 1, L079: 1, L077: 1, L0799: 1, L0687: 1, L0599: 1, H0136: 1, S0242: 1, S0194: 1, S0276: 1, H0543: 1, S0424: 1 and S0452: 1.					H0135: 2, S0046: 1, S0222: 1, H0052: 1, H0672: 1, S0328: 1 and		
							Gly-16 to Lys-22, Trp-82 to Lys-87.
	4279	4280	4281	4282	4283	4284	4285
	47 - 754	338 - 778	2 - 214	3 - 497	1 - 441	2-415	3 - 350
	2104	2105	2106	2107	2108	2109	2110
	HVVCD90 HVVCD90R	HVVCE63 HVVCE63R	HVVCE65R	HVVCF38R	HVVCG29R	HVVCG31R	HVVCG46 HVVCG46R
	HVVCD90	HVVCE63	HVVCE65	HVVCF38	HVVCG29	HVVCG31	HVVCG46

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Arg-48 to Asn-57, AR089: 107, AR096: Lys-77 to Glu-83, 52, AR104: 30, AR060: Phe-106 to Asp-112, 27, AR061: 14, AR055: 13		·				H0585: 4, H0457: 4,	L0809: 4, L0747: 4,	S0442: 3, H0492: 3,	H0427: 3, H0318: 3,	L0666: 3, H0690: 3,	H0686: 2, H0662: 2,	S0358: 2, S0408: 2,	H0150: 2, H0135: 2,	H0063: 2, H0494: 2,	L0761: 2, L0764: 2,	L0773: 2, L0662: 2,	L0378: 2, L0665: 2,	H0670: 2, S0406: 2,	L0744: 2, L0749: 2,	L0601: 2, H0170: 1,
Arg-48 to Asn-57, Lys-77 to Glu-83, 5 Phe-106 to Asp-112.2	Glu-81 to Gly-91, Arg-107 to Asn-116.	Asp-10 to Lys-15, Lys-29 to Asp-34.		Asp-1 to Thr-14, Ala-76 to Thr-88,	Asp-101 to Lys-115, Ala-128 to Gln-136, Pro-141 to Gln-151.								-			-				
4286		4288	4289	4290		1201	1671													
1 - 405	3 - 350	3 - 119	44 - 460	3 - 647		1 251	1-331													
2111	2112	2113	2114	2115			2110													
HVVCG93 HVVCG93R	HVVCH28R	HVVCI28R	HVVCI50R	HVVCI85R	-		HVVCJ51K											-	•	
HVVCG93	HVVCH28	HVVCI28	HVVCTS0	HVVCI85			HVVCJ51													

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H0685: 1, H0661: 1,	H0402: 1, S0348: 1,	S0354: 1, S0376: 1,	S0360: 1, S0278: 1,	H0586: 1, H0333: 1,	H0642: 1, L0622: 1,	S0280: 1, H0599: 1,	H0253: 1, H0231: 1,	H0545: 1, H0373: 1,	H0266: 1, H0553: 1,	S0036: 1, H0551: 1,	H0116: 1, H0264: 1,	H0059: 1, S0352: 1,	S0440: 1, H0646: 1,	L0640: 1, L0639: 1,	L0630: 1, L0772: 1,	L0373: 1, L0646: 1,	 L0374: 1, L0644: 1,	L0645: 1, L0794: 1,	-	<u> </u>	L0793: 1, L0532: 1,	L0664: 1, H0520: 1,	H0672: 1, S0380: 1,	H0696: 1, H0478: 1,		L0750: 1, L0759: 1,	_	H0422: 1.
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L0751: 3, L0757: 3, L0662: 2, H0686: 1, H0266: 1, L0055: 1, L0763: 1, L0764: 1, L0768: 1, L0805: 1, L0653: 1, L0666: 1, H0690: 1, H0672: 1, L0777: 1, L0758: 1 and S0276: 1.							H0521: 182, H0522: 34, S0002: 30, S0278: 27, H0638: 23, S0344:
		Pro-11 to Lys-31, Ser-36 to Gln-44, Thr-53 to Gly-58, Ala-74 to Arg-87, Pro-105 to His-113.	Gly-8 to Lys-22, Phe-29 to Trp-34, Val-95 to Cys-100, Thr-147 to Gln-161, Gln-164 to Lys-170, Glu-179 to Leu-185, Pro-194 to Gly-199.	Trp-36 to Thr-43, Gly-51 to Thr-70, Tyr-87 to Glu-97.			
4292	4293	4294	4295	4296	4297	4298	4299
1 - 147	3 - 320	1 - 339	2-775	3 - 293	3 - 347	35 - 466	96 - 332
2117	2118	2119	2120	2121	2122	2123	2124
HVVCK59 HVVCK59R	HVVCK78R	HVVCL52R	HVVCL73R	HVVCM67 HVVCM67R	HVVCM84R	HVVCN20R	HVVCN29 HVVCN29R
HVVCK59	HVVCK78		HVVCL73	HVVCM67	HWWCM84	THAN CANDO	HVVCN29

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			Gln-61 to Pro-68,	Gln-103 to Ala-108,	Tyr-111 to Thr-117.	Val-28 to Ser-33.	Lys-54 to Ala-67,	Arg-77 to Thr-86.		Ala-35 to Ala-41,	Pro-46 to Gly-54,	Pro-67 to Gly-87.	Ser-23 to Lys-30.	Glu-74 to Arg-79,	Glu-117 to Arg-134.	Arg-30 to Phe-36.			Arg-11 to Pro-17,	Uis 1 to A on 0	to-lie of 1-sin	Arg-10 to Gly-16,	Tyr-50 to Gly-56.	
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	HVVCN54	HVVCN66	HVVC013			HVVC016	HVVCO87		HVVCP41	HVVCP88			HVVCQ05	HVVCQ49		HVVCQ70	HVVCQ93	HVVCS28	HVVCS32	111111111111111111111111111111111111111		_		

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	Gln-8 to Lys-13, Glu-32 to Arg-56, Ser-60 to Thr-69.
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	Glu-1 to Gln-10,	Pro-12 to Ser-18, Pro-25 to Pro-30,	Met-22 to Asp-27, Cys-43 to Trp-93, Asp-79 to Trp-93, Glu-124 to Glu-136.
-	4317 G	4318 P	4319 V V V
	2-310	2 - 244	3-410
	2142	2143	2144
	HVVCV41R	HVVCV68 HVVCV68R	HVVCW75R
	HVVCV41 HVVCV41R	HVVCV68	HVVCW75

0601: 1 and	H0052: 15, H0585: 12,	L0439: 11,	L0754: 9,	0748: 8,	0666: 7,	0742: 7,	0556: 6,	0494: 6,	0751: 6,	0253: 5,	0803: 5,	.0657: 4,	0408: 4,	0278: 4,	0049: 4,	10623: 4,	0764: 4,	0805: 4,	0744: 4,	0040: 3,	0005: 3,	10333: 3,	10599: 3,	10242: 3,	I0038: 3,	10412: 3,	0769: 3,	0774: 3,
L0592: 1, L0601: 1 and S0424: 1.	Γ	L0740: 12, L0439: 11,	L0758: 10, L0754: 9	L5286: 8, L0748: 8,	H0141: 7, L0666: 7	H0521: 7, L0742: 7	L0749: 7, H0556: 6,	S0222: 6, H0494: 6,	L0743: 6, L0751: 6,	L0757: 6, H0253: 5,	T0010: 5, L0803: 5,	L0731: 5, H0657: 4	S0360: 4, S0408: 4,	S0046: 4, S0278: 4,	H0618: 4, S0049: 4	H0617: 4, H0623: 4	S0038: 4, L0764: 4,	L0766: 4, L0805: 4,	L0809: 4, L0744: 4,	L0746: 4, S0040: 3,	S0212: 3, L0005: 3	H0441: 3, H0333:	H0013: 3, H0599: 3	H0009: 3, H0242: 3,	H0644: 3, H0038:	H0551: 3, H0412: 3,	T0042: 3, L0769: 3,	L0804: 3, L0774: 3,
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<u>其民政政政立 其中七十十十二日 </u>	Arg-9 to Ile-31, Gln-38 to Lys-43, Gln-87 to Ala-92,
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Ser-116 to Pro-128.	Ser-8 to Ser-14,	Asn-68 to Tyr-74,	His-80 to Val-91,	Glu-141 to Phe-147,	Ala-159 to Ser-170,	Asp-184 to Gly-199.	Pro-60 to Gly-72,	Gln-89 to Ser-94,	Pro-121 to Pro-132,	Pro-146 to Pro-166.	Lys-30 to Glu-36,	Lys-55 to Tyr-68,	Ile-82 to Asp-87,	Trp-115 to Ser-125,	Pro-139 to Pro-146.	Ala-17 to Lys-23,	Leu-25 to Ala-35,	Gln-71 to Ala-76,	Tyr-79 to His-85,	Pro-112 to Gly-120,	Gly-149 to Tyr-165,	Lys-175 to His-181.		Glu-179 to Ala-184.				
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	2-775						2 - 697				1 - 588					3 - 620							139 - 276	2 - 598	423 - 542	139 - 291		
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	HVVCY29R						HVVCYSSR				HVVCY60 HVVCY60R					HVVCY62 HVVCY62R							HVVCY75R	HVVCY77 HVVCY77R 2	HVVCZ18R	HVVCZ50R		
	HVVCY29						HVVCY55				HVVCY 60					HVVCY62							HVVCY75	HVVCY77	HVVCZ18	HVVCZ50		

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3005	H0411: 5, H0486: 4	S0003: 4, S0210: 4,	H0672: 4, L0731: 4,	S0192: 4, S0418: 3,	30046: 3, H0169: 3,	S004	H066	S037	H063	H05	H05	H00	H05	H04	1.05	90H	30028: 2, H0506: 2	H0170: 1, H0171:	, H0661:	10639: 1, H0574: 1	,10022:	H0108: 1, H0042:	H0004: 1, H0044:	H0355: 1, H0510:	H0266: 1, S0318: 1	S0316: 1, S0022: 1	, HO	Ä,	Ĕ,
5.5	1:5,	3.4.5	2: 4,	2: 4, 5	6:3,1	9:3	1:2,	8:2	2:2	8: 2,	16: 2,	12: 2,	9.7	58: 2,	59. 2,	9.2	28: 2,	70: 1	H0295: 1,	39: 1	H0599: 1, I	08:1	1.40	55: 1	66: 1	16:1	128: 1	522: 1	188: 1
S0212: 5, S0045: 5,	1041	3000	H067	S019.	S004	1064	H067	S035	S013	500H	H054	H024	S025	H02	106	108	2002	H01	H02	H06	H05	H01	0H	H03	H02	S03	H03	HOH	Ę H H O H
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	HVVDC09R
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10713: 1,	0661: 1,	0360: 1,	0580: 1,	0619: 1,	6022: 1,	, H0013: 1,	10097: 1,	0110: 1,	.0471: 1,	10687: 1,	0214: 1,	0023: 1,	10551: 1,	10494: 1,	0352: 1,	10633: 1,	10538: 1,	0426: 1,	.0500: 1,	L0637: 1,	.0764: 1,	.0774: 1,	L0375: 1,	L0805: 1,	.0659: 1,	.0666: 1,	.0665: 1,	10148: 1,	H0519: 1,
H0222: 1, H0713: 1	L0420: 1, H0661:	H0663: 1, S	S0410: 1, H0580: 1	S0007: 1, H	H0351: 1, S	H0587: 1, H	H0427: 1, H	H0052: 1, T0110: 1	H0231: 1, L	H0266: 1, H0687: 1	S0250: 1, S0214: 1	H0428: 1, T0023: 1,	H0628: 1, F	H0412: 1, H0494:	H0366: 1, S0352:	S0440: 1, H0633:	H0646: 1, H0538:	S0422: 1, S0426:	L0369: 1, L0500: 1	L0770: 1, L		L0381: 1, I	L0775: 1, I	L0806: 1, I	L0654: 1, I	L0783: 1, I	L0664: 1, L0665: 1,	S0374: 1, S	H0520: 1, H0519:
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	4331	4332	4333	4334																		
	2 - 580	3 - 545	3 - 446	1 - 390																		
	2156	2157	2158	2159																		
	HVVDH44 HVVDH44R		HVVDM31R	HVVDM45R							•											
·	HVVDH44	HVVDM23	HVVDM31	HVVDM45																		

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H0544: 1, H0046: 1, H0009: 1, H0023: 1, N0007: 1, T0010: 1, H0188: 1, H0687: 1, S0364: 1, L0455: 1, S0366: 1, H0050: 1, H00412: 1, H0059: 1, H0641: 1, S0142: 1, S0002: 1, H057: 1, L077: 1, L077: 1, L0775: 1, L056: 1, L0655: 1, L0352: 1, S0008: 1, H0435: 1, H0672: 1, H0658: 1, H0576: 1, L0779: 1, L0605: 1, L0593: 1, S0276: 1, L0593: 1, S0276: 1, H0423: 1 and H0542: 1, H0423: 1 and H0542: 1, H0423: 1 and		L0766: 9, L0665: 7, L0659: 6, L0761: 5, L0662: 4, L0776: 4,
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S0418: 3, S0007: 3,	H0559: 3, H0457: 3,	H0617: 3, L0800: 3,	L0751: 3, L0750: 3,	H0309: 2, L0769: 2,	L0667: 2, L0794: 2,	L0664: 2, L0756: 2,	L0779: 2, L0777: 2,	L0758: 2, H0543: 2,	H0556: 1, S0402: 1,	S0114: 1, H0650: 1,	H0656: 1, S0116: 1,	H0484: 1, H0254: 1,	H0255: 1, S0420: 1,	S0354: 1, S0358: 1,	H0676: 1, L0717: 1,	S0222: 1, H0614: 1,	L0622: 1, H0486: 1,	H0069: 1, H0253: 1,	H0581: 1, H0024: 1,	H0051: 1, H0100: 1,	H0494: 1, H0625: 1,	H0561: 1, S0144: 1,	L0374: 1, L0771: 1,	L0773: 1, L0768: 1,	L0387: 1, L0774: 1,	L0653: 1, L0657: 1,	L0384: 1, L0789: 1,	L0790: 1, L0792: 1,	H0684: 1, H0659: 1,
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	Asp-50 to Gly-55.		Lys-24 to Glu-34,	Fro-4 / to Arg-56, Tyr-77 to Pro-90.	Lys-12 to Thr-18,	Glu-20 to Lys-26,	Lys-83 to Lys-90.	•																
	4337	4338	4339		4340			•	•															
	2 - 685	1 - 594	40 - 354		106 - 423					-										,				
	2162	2163	2164		2165													,						
	HVVDN29R	HVVDN77 HVVDN77R	HVVDP70 HVVDP70R		HVVDO46 HVVDO46R	y 1																		
	HVVDN29	HVVDN77	HVVDP70		HVVDO46	2									-									

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L0761: 2. L0773: 2.	0649: 2, L0774: 2,		10435: 2, L0749: 2,	.0731: 2, L0588: 2,	.0608: 2, S0026: 2,	HO543: 2, H0294: 1,	10650: 1, H0341: 1,	S0354: 1, S0376: 1,	10580: 1, H0339: 1,	10208: 1, S0132: 1,	10640: 1, H0369: 1,	10438: 1, H0574: 1,	10486: 1, H0101: 1,	H0427: 1, H0097: 1,	•	10230: 1, H0052: 1,	.0040: 1, H0544: 1,	10545: 1, H0086: 1,	10009: 1, S0024: 1,	10510: 1, S0250: 1,	10428: 1, H0166: 1,	H0102: 1, H0560: 1,	S0294: 1, H0509: 1,	S0150: 1, H0633: 1,	10647: 1, L0631: 1,	.0646: 1, L.0764: 1,	.0804: 1, L0806: 1,	.0606: 1, L0657: 1,	.0659: 1, L0636: 1,
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L0783: 1, L0382: 1, L0789: 1, L0664: 1, L0665: 1, H0593: 1, H0690: 1, H0682: 1, H0672: 1, H0648: 1, H0672: 1, H0539: 1, H0672: 1, L0609: 1, L0744: 1, L0748: 1, L0747: 1, L0755: 1, H0445: 1, H0343: 1, L0581: 1, H0653: 1 and H0352: 1.	H0521: 7, H0052: 6, H0622: 5, L0646: 5, H0617: 4, L0662: 4, L0659: 4, H0556: 3, H0341: 3, H0484: 3, S0358: 3, H0580: 3, H0551: 3, L0664: 3, H0670: 3, S0328: 3, H0670: 3, S0328: 3, H0583: 2, S0466: 2, S0376: 2, S0046: 2, S0132: 2, H0438: 2, H0586: 2, H0486: 2, H0618: 2, S0474: 2, S0049: 2, H0545: 2,
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	HVVDQ4

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H0268: 2, H0494: 2,	H0509: 2, H0647: 2,	.5565: 2, L0645: 2,	0764: 2, L0666: 2,	,0665: 2, H0520: 2,	H0689: 2, H0658: 2,	S0152: 2, S0406: 2,	S3014: 2, L0439: 2,	.0751: 2, L0754: 2,	0750: 2, L0758: 2,	S0434: 2, L0596: 2,	0581: 2, L0595: 2,	.0601: 2, S6024: 1,	10294: 1, H0657: 1,	H0656: 1, L0415: 1,	H0669: 1, H0671: 1,	56: 1, S0354: 1,	S0360: 1, S0410: 1,	H0340: 1, S0476: 1,	S0278: 1, H0549: 1,	50: 1, H0370: 1,	H0587: 1, H0497: 1,	10643: 1, H0257: 1,	H0318: 1, H0457: 1,	H0041: 1, H0123: 1,	H0024: 1, H0071: 1,	H0375: 1, H0290: 1,	H0292: 1, H0428: 1,	.0142: 1, S0036: 1,	H0063: 1, H0087: 1,
H02	H05	LSS	101	106	90H	801	S30	101	101	S04.	105	907	H02	90H	90H	803	203	H03	205	HOS	H05	90H	H03	H00	00H	E0H	H02	ioi	00H
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H0264: 1, H0059: 1, H0100: 1, S0448: 1, S0450: 1, S0448: 1, S0052: 1, L0762: 1, L0640: 1, L0639: 1, L0643: 1, L0771: 1, L0643: 1, L0771: 1, L0773: 1, L0771: 1, L0773: 1, L0774: 1, L0775: 1, L0774: 1, L0775: 1, L0778: 1, H0806: 1, L0774: 1, L0776: 1, L0778: 1, H0693: 1, H0672: 1, S0330: 1, H0696: 1, H0519: 1, H0478: 1, S0028: 1, L0747: 1, L0749: 1, L0731: 1, S0436: 1, L0731: 1, S0436: 1, L0743: 1,	H0423: 1 and H0352: 1	AR104: 17, AR089: 10, AR096: 10, AR061: 8, AR060: 4, AR055:
	Paris I	1
	4342	4343
	1 - 732	2 - 730
	2167	1
	HVVDS12R	HVVDS21R
	HWD812	HVVDS21

4	S0412: 22, L0662: 19,	LU///: 14, S0010: 11,	S0222: 9, S6028: 9,	L0750: 9, T0010: 8,	L0659: 8, L0747: 8,	L0756: 8, L0663: 7,	L0439: 7, H0051: 6,	L0518: 6, L0754: 6,	L0752: 6, S0280: 5,	H0575: 5, S0358: 4,	SQ346: 4, L0753: 4,	H0170: 3, H0662: 3,	S0360: 3, H0427: 3,	L0471: 3, H0373: 3,	L0638: 3, L0637: 3,	L0764: 3, L0774: 3,	L0809: 3, L0666: 3,	S0310: 3, H0672: 3,	H0696: 3, L0731: 3,	H0506: 3, H0713: 2,	S0408: 2, H0411: 2,	H0455: 2, H0574: 2,	S0414: 2, H0486: 2,	H0036: 2, S0049: 2,	H0052: 2, H0327: 2,	H0687: 2, H0428: 2,	H0553: 2, H0644: 2,	H0038: 2, L0520: 2,
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L0762: 2, L0770: 2, L0653: 2, L0517: 2, S0044: 2, L0442: 1, H0583: 1, L0443: 1, \$0110: 1, S0282: 1, S0400: 1, H0661: 1, S0348: 1, S0356: 1,	S0444: 1, H0329: 1, S0007: 1, H0619: 1, S6026: 1, S0300: 1, L0717: 1, S6022: 1, H0550: 1, H0592: 1, H0587: 1, H0599: 1, H0042: 1, H0399: 1,	H0545: 1, H0563: 1, H0564: 1, H0123: 1, H0019: 1, S0050: 1, L0163: 1, S0388: 1, H0356: 1, H0328: 1, H0688: 1, H0039: 1, T0023: 1, H0031: 1,	H0111: 1, H0169: 1, S0364: 1, L0455: 1, H0135: 1, H0163: 1, T0067: 1, T0069: 1, T0004: 1, H0100: 1, S0112: 1, L0370: 1, L0598: 1, L0769: 1, L0630: 1, L0800: 1,
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L0648: 1, L0363: 1, L0768: 1, L0649: 1, L0803: 1, L0775: 1, L0375: 1, L0651: 1, L0805: 1, L076: 1, L0805: 1, L0776: 1, L0805: 1, L0776: 1, L0635: 1, L0778: 1, L0664: 1, H0691: 1, H0693: 1, H0691: 1, H0693: 1, L078: 1, H0684: 1, S0174: 1, H0684: 1, S0174: 1, L0743: 1, L0748: 1, L0779: 1, L0749: 1, L0779: 1, L0749: 1, L0779: 1, L0769: 1, L0689: 1, S0434: 1, L0604: 1, L0366: 1, S0106: 1 and S0021: 1.	
	Thr-35 to Asn-41
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	L0748: 11, L0758: 10, L0747: 8, L0439: 7,	H0144: 6, L0805: 5,	L0662: 4, H0341: 3,	H0052: 3, L0794: 3,	L0803: 3, L0809: 3,	H0672: 3, L0741: 3,	L0752: 3, H0619: 2,	S0222: 2, H0013: 2,	H0014: 2, H0266: 2,	H0553: 2, L0455: 2,	H0591: 2, L0770: 2,	L0646: 2, L0764: 2,	L0666: 2, L0665: 2,	H0547: 2, S0126: 2,	L0596: 2, L0589: 2	L0605: 2, H0543: 2,	H0423; 2, H0624: 1,	H0556: 1, H0686: 1,	S0040: 1, H0717:	H0671: 1, S0476:	L0717: 1, H0411:	H0586: 1, H0333:	H0574: 1, L0021:	H0575: 1, H0618:	S0010: 1, H0318:	H0544: 1, H0009: 1	H0050: 1, L0471: 1,	L0163: 1, S6028:
	Glu-11 to Arg-17,		Ser-116 to Ser-125.	_			•																					
4346	4347																											
109 - 471	2 - 460											ì																
2171	2172						,																		•			
HVVDT29R	HVVDT44 HVVDT44R		,																									
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S0022: 1, S0214: 1, H0628: 1, L0456: 1, S0366: 1, S0366: 1, H0039: 1, H0039: 1, H0040: 1, H0059: 1, S0112: 1, H0647: 1, H0647: 1, H0647: 1, H0647: 1, L0638: 1, L0768: 1, L0639: 1, L0768: 1, L0649: 1, L0774: 1, L0649: 1, L0647: 1, L0664: 1, S0053: 1, L0664: 1, S0053: 1, H0659: 1, L0744: 1, L0754: 1, L0744: 1, L0754: 1, L0744: 1, L0754: 1, L0608: 1, L0608: 1, L0608: 1, L0608: 1, H0665: 1, H06422: 1, H06422: 1	H0672: 2 and S6028: 1.
	Asn-1 to Leu-6.
	4348
	328 - 600
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	HVVDU49

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	Ser-1 to Pro-9.							Lys-1 to Asp-8, Lys-41 to Gln-50.
4349		4351	4352	4353	4354	4355		4356
3-266	2 - 502	294 - 455	14 - 511	2 - 391	11 - 178	167 - 343		3 - 197
2174	_	+-	21.77	2178	2179	2180		2181
dcoluciaring	HVVDW02 HVVDW02K	HVVDX40R	HVVDX90 HVVDX90R	HWHHD07R	HWI ACSIR	HWLJR51 HWLJR51R		HWLME48 HWLME48R
	HVVDW02	HVVDW01	OBXUVVII	מממאאות	HWIT A CS1	HWLJR51		HWLME48

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	Cys-6 to Ser-12, Gln-27 to Glu-32,	Arg-52 to Arg-67, Asp-140 to Asp-	Thr-26 to Ile-34.
4357	4358		4359
209 - 373	1 - 435		43 - 210
2182	2183		2184
HWLVY65R	HWMIB35R		HWNF025R
HWLVY65	HWMIB35		HWNFO25 HWNFO25R
	HWLVY65R	2182 209 - 373 4357 2183 1 - 435 4358	2182 209 - 373 4357 2183 1 - 435 4358

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	1 - 300
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¥	HWWEF90 HWWEF90R
	HWWEF90 HWWE

[0040] The first column in Table 1 provides a unique "Clone ID NO:Z" for a cDNA clone related to each contig sequence disclosed in Table 1. This clone ID references the cDNA clone which contains at least the 5' most sequence of the assembled contig, and at least a portion of SEQ ID NO:X was determined by directly sequencing the referenced clone. The reference clone may have more sequence than described in the sequence listing or the clone may have less. In the vast majority of cases, however, the clone is believed to encode a full-length polypeptide. In the case where a clone is not full-length, a full-length cDNA can be obtained by methods known in the art and/or as described elsewhere herein.

The second column in Table 1 provides a unique "Contig ID" identification for each contig sequence. The third column provides the "SEQ ID NO:X" identifier for each of the ovarian associated contig polynucleotide sequences disclosed in Table 1. The fourth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence "SEQ ID NO:X" that delineate the preferred open reading frame (ORF) shown in the sequence listing and referenced in Table 1, column 5, as SEQ ID NO:Y. Where the nucleotide position number "To" is lower than the nucleotide position number "From", the preferred ORF is the reverse complement of the referenced polynucleotide sequence.

[0042] The fifth column in Table 1 provides the corresponding SEQ ID NO:Y for the polypeptide sequence encoded by the preferred ORF delineated in column 4. In one embodiment, the invention provides an amino acid sequence comprising, or alternatively consisting of, a polypeptide encoded by the portion of SEQ ID NO:X delineated by "ORF (From-To)". Also provided are polynucleotides encoding such amino acid sequences and the complementary strand thereto.

[0043] Column 6 in Table 1 lists residues comprising epitopes contained in the polypeptides encoded by the preferred ORF (SEQ ID NO:Y), as predicted using the algorithm of Jameson and Wolf, (1988) Comp. Appl. Biosci. 4:181-186. The Jameson-Wolf antigenic analysis was performed using the computer program PROTEAN (Version 3.11 for the Power MacIntosh, DNASTAR, Inc., 1228 South Park Street Madison, WI). In specific embodiments, polypeptides of the invention comprise, or alternatively consist of, at least one, two, three, four, five or more of the predicted epitopes as described in Table 1. It

will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly.

Column 7 in Table 1 provides an expression profile and library code: count [0044] for each of the contig sequences (SEQ ID NO:X) disclosed in Table 1, which can routinely be combined with the information provided in Table 4 and used to determine the normal or diseased tissues, cells, and/or cell line libraries which predominantly express the polynucleotides of the invention. The first number in column 7 (preceding the colon), represents the tissue/cell source identifier code corresponding to the code and description provided in Table 4. For those identifier codes in which the first two letters are not "AR", the second number in column 7 (following the colon) represents the number of times a sequence corresponding to the reference polynucleotide sequence was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array technology. Utilizing this technology, cDNAs were amplified by PCR and then transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array. cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of ³³P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression. The sequences disclosed herein have been determined to be

predominantly expressed in ovarian tissues, including normal and diseased ovarian tissues (See Table 1, column 7 and Table 4).

[0045] Column 8 in Table 1 provides a chromosomal map location for certain polynucleotides of the invention. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Each sequence in the UniGene database is assigned to a "cluster"; all of the ESTs, cDNAs, and STSs in a cluster are believed to be derived from a single gene. Chromosomal mapping data is often available for one or more sequence(s) in a UniGene cluster; this data (if consistent) is then applied to the cluster as a whole. Thus, it is possible to infer the chromosomal location of a new polynucleotide sequence by determining its identity with a mapped UniGene cluster.

[0046] A modified version of the computer program BLASTN (Altshul et al., J. Mol. Biol. 215:403-410 (1990), and Gish et al., Nat. Genet. 3:266-272 (1993)) was used to search the UniGene database for EST or cDNA sequences that contain exact or near-exact matches to a polynucleotide sequence of the invention (the 'Query'). A sequence from the UniGene database (the 'Subject') was said to be an exact match if it contained a segment of 50 nucleotides in length such that 48 of those nucleotides were in the same order as found in the Query sequence. If all of the matches that met this criteria were in the same UniGene cluster, and mapping data was available for this cluster, it is indicated in Table 1 under the heading "Cytologic Band". Where a cluster had been further localized to a distinct cytologic band, that band is disclosed; where no banding information was available, but the gene had been localized to a single chromosome, the chromosome is disclosed.

Once a presumptive chromosomal location was determined for a polynucleotide of the invention, an associated disease locus was identified by comparison with a database of diseases which have been experimentally associated with genetic loci. The database used was the Morbid Map, derived from OMIMTM (supra). If the putative chromosomal location of a polynucleotide of the invention (Query sequence) was associated with a disease in the Morbid Map database, an OMIM reference identification number was noted in column 9, Table 1, labeled "OMIM Disease Reference(s)". Table 5 is a key to the OMIM reference identification numbers (column 1), and provides a description of the associated disease in Column 2.

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Clone ID NO:Z	Contig ID:	SEQ ID	Analysis Method	PFam/NR Description	Pram/NK Accession	Percent Identity	From	l I
		×			11 201304 4 01:31	28%	319	167
HOVCD34	396327	111	blastx.2	(AK000844) unnamed protein product [Homo	db BAA91390.1			-
				sapiens	-:- A 44805 A 44805	74%	3	86
HEBGD58	498281	12	blastx.2	eggshell protein - fluke (Schistosoma				
				haematobium)				
				(subclone I	-LIA A A 88848 11	100%	06	947
HETCD42	533532	13	blastx.2	actin capping protein alpha subunit [Homo	goldzenog			
		_		sapiens	11 3001004 1 211 11	%69	873	9/9
HTXKC18	535854	14	blastx.2	(AK000496) unnamed protein product [Homo	dbj BAA91203.1 	~ 		
				sapiens	1104 4 51280 11	91%	150	1283
HLDRK20	553765	15	blastx.2	complement factor B	emb[CAA51389.1]			
	一	14	hlacty 2	NBR2 PROTEIN	sp 015453 NBR2_H	100%	138	7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7
HZMBD33	5284/4	2		(NEXT TO BRCA1	UMAN			
				GENE 2 PROTEIN).	10000	7080		3 515
19XBV9U	558708	17	blastx.2	N-cadherin [Homo	gb AAAU3236.1			
TOTAL CIT				sapiens	114 4 E02026 11 4 E1	%19	1063	3 905
HELHC03	562745	138	blastx.2	(AF118082) PRO1902	gb AAF22020.1 54.1 18094 21			
				Homo sapiens	emblCA A33261.11	75%	6 120	0 347
UCENAD70	585385	19	blastx.2	precursor polypepude	- I consolored			 -
1 TAT. TOG —	_							

	989	687	746		466	498	393		434	434	431	434	431	431	431	431	431	431	431	431	431	431	431	434	431	431	431
	522	19	681		7	463	133		117	192	189	12	192	195	192	156	189	156	195	195	192	15	192	195	192	15	105
**	%E9	%66	%06		100%	91%	100%		100%	20%	51%	40%	51%	52%	23%	47%	24%	46%	\$2%	47%	48%	38%	48%	20%	46%	37%	7007
	gb[AAA88038.1	gb[AAA59491.1]	- -		gb AAD34121.1 AF1	51884 1	gb AAD02337.1		gb AAC33512.1					-													
(AA -21 to 782) [Homo sapiens]	unknown protein [Homo sapiens]	leukocyte adhesion	glycoprotein precursor	Homo sapiens	(AF151884) CGI-126	protein [Homo sapiens]	(AF044671) MIM46	[Homo sapiens]	(AF019406) collagen	type IX alpha 2 chain	Homo sapiens	1								,							
	blastx.2	blastx.2			blastx.2		blastx.2		blastx.2								•	-	•								
	70	21	· ·		77		23		. 24															_			
-	585675	588869			638220		654868		704405																	.•	
	HSKNZ25	HDPFK39			HL1AB07		HHGAS83		HTSGU37												•						

431	431	431	431	431	431	434	425	461	434	413	395	365	431	347	502	490	481	472	481	481	481	481	481	205	487	457	432	218	133
195	201	24	189	189	180	195	192	195	195	195	45	192	195	192	395	395	383	401	401	383	395	383	383	380	422	419	331	430	249
48%	%05	37%	48%	48%	44%	46%	43%	40%	43%	48%	34%	44%	36%	47%	47%	26%	44%	52%	46%	38%	43%	40%	38%	33%	47%	61%	29%	52%	37%
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240	47	501	1	402					101	58	14	619	61	77	969	22	90	122	34	12	20		
38%	20%	32%	64%	87%					%56	95%	21%	35%	32%	38%	73%	77%	63%	%68	70%	81%	%98		
			emb CAB42187.1	emb CAA57432.1			-		emb CAA90511.1						gb[AAA88038.1]			gb AAC78561.1		emb CAA45263.1			
			unnamed protein product [unidentified]	fusion protein:	ubiquitin (bases	43_513); ribosomal	protein S27a (bases	217_532) [Kattus norvegicus]	tyrosine kinase	activator protein 1	(TKA-1) [Homo	sapiens			unknown protein	Homo sapiens		(AF046001) zinc finger	transcription factor [Homo sapiens]	cytochrome oxidase	subunit III [Phoca vitulina]		
			blastx.2	blastx.2					blastx.2						blastx.2			blastx.2	,	blastx.2			
			25	76					27						28			59		30			
			705692	711500					732342						745343			746416		762806			,
	:		H2LAN34	HPMBZ40					HLYEP52						HTTEC47			HOFMO90		HCHIND34			

HAMGI86	785328	31	blastx.2	unnamed protein	emb CAB42212.1	91%	424 611	609 667
HLWCN67	794213	32	blastx.2	(AK014485) putative	dbj BAB29387.1	%08	386	631
HOFAC09	806819	33	blastx.2	acidic ribosomal phosphoprotein (P0)	gb AAA36470.1	87% 52%	42 5	812
HRDEL61	824886	34	blastx.2	X-linked retinopathy protein [C-terminal, clone XEH.8c] [human, Peptide Partial, 100 aa]	gb AAB26149.1	67% 45%	808 644	632 576
HDPOR60	828176	35	blastx.2	(AF151799) CGI-40	gb AAD34036.1 AF1 51799 1	96% 82%	∞ ∞	1867 1867
HPRTS71	828574	36	blastx.2	rTSbeta [Homo	emb CAA61761.1	100%	45	458
HOHBI90	828862	37	blastx.2	tyrosine kinase receptor [Homo sapiens]	gb AAA61243.1	90% 63% 34%	193 159 268 375	417 281 396 419
HOHAL47	828872	38	blastx.2	pericentriol material 1	gb AAA60120.1	%88	328	879
HYASE58	829298	39	blastx.2	O-6-methylguanine- DNA methyltransferase (Homo sapiens]	gb AAA59594.1	%66	74	694
HKAAH95	829958	40	blastx.2	ribosomal protein L22	emb CAA55204.1	70%	41	418

		Ŀ		[Rattus norvegicus]				
HTTIQ02	829981	41	blastx.2	activating transcription	gb AAA20506.1	%96	44	391
				factor 3 [Homo				
				sapiens				
HWACG9	830195	42	blastx.2	heat shock protein 90,	gb AAB23369.1	82%	119	610
-				hsp90 [rats, brain,		93%	88	132
				Peptide, 724 aa] [Rattus	٠	37%	9	170
				sp.]		%9/	573	611
HUFBX52	830497	43	blastx.2	tenascin X [Homo	gb AAB47488.1 AAB	%66	2	1525
				sapiens]	47488	100%	1525	1785
						767	70	1255
						79%	20	1114
						30%	20	1033
						27%	17	991
						79%	70	886
						78%	17	934
						25%	70	1096
			,			27%	∞	868
						78%	7	. 973
						78%	14	856
						27%	70	892
			,		-	78%	17	712
				,		767	70	712
					. •	24%	365	1150
	. ,					24%	70	973
						32%	413	946
	•					78%	11	200
						24%	7	991
	. •		÷	:		30%	20	289
						32%	17	589

712	601	856	1111	1012	295	1096	595	1135	1045	1096	1231	1141	1111	1153	1309	1135	1117	1021	604	1690	1096	1684	1135	1309	1684	877	331	1045	1684
2	70	7	716	176	176	437	6	521	437	434	488	860	557	860	557	608	641	437	437	1487	557	1487	788	608	1487	638	35	455	1487
73%	27%	27%	31%	25%	30%	25%	73%	25%	23%	23%	70%	25%	22%	25%	24%	28%	20%	27%	36%	36%	27%	78%	23%	22%	30%	33%	75%	23%	28%
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1684	825	1684	1690	1684	1690	501	1190	1150	1142	404	498	1663	1181	1181	1027	1436	1118	257	255	1083	1146	480	1175	315	357	181	266	1305
1487	622	1487	1556	1487	1487	316	1110	962	1101	267	52	1607	1089	1089	650	1398	1089	189	118	626	1024	361	1050	46	175	143	134	1018
27%	30%	25%	37%	78%	23%	27%	48%	30%	64%	32%	76%	25%	32%	32%	22%	23%	%09	41%	75%	37%	29%	35%	34%	%86	100%	95%	100%	92%
											-											•		gb AAA35886.1	emb CAA28169.1		gb AAD09765.1	
																							•	2A9 peptide [Homo sapiens]	precursor polypeptide	[Homo sapiens]	(AF096304) putative	sterol reductase SR-1
			-																			-		blastx.2	blastx.2		blastx.2	
																	·							44	45		46	
										-														831453	832454		833088	
					,								-										•	HWLJE49	HLOBT44	,	HSLGG58	

1029	480 170	353		1096	834	301	1102	172		382	\$00	COO	266	
926	157 48	123		404	1121	591 281	1188	11		11	tot) 	406	485 333
61%	95%	100%		100%	29%	34%	41%	100%		48%		93%	55%	71% 59%
	dbj BAA21510.1	gb AAB61158.1		gb AAA60239.1	11 4 4 777 810 11	gb AAC/2010:1		gb[AAC52117.1]		emblCA A90017.1		gb AAB34132.1	GHIA 4 F78916.11AF1	61356_1
	rig-analog g protein	Sus scrofa cysteine-rich intestinal protein [Homo sapiens]		winone oxidoreductase	[Homo sapiens]	(AF081114) ORF2 [Mus musculus	domesticus]	1. Landa like motor	kinesin-like inoka protein KIF1C [Homo	sapiens	plasmolipin [Kauus norveoicus]	P26s4 [Cricetinae gen.	sp.]	(AF161356) H3FC093 [Homo sapiens]
	blastx.2 (blastx.2 0			blastx.2	blastx.2			blastx.2		blastx.2	blastx.2		blastx.2
	47	48	·		49	50			51		25	53		54
	840756	840862			841088	843485			844534		847355	047647) to / to .	847821
	нснвозз 8	HDTMK30			HDPFX64	HODFG71			HCE3165		HSAVH65	100000	HBAF141	HBMCM3

						63%	534	502
HACBO42	849064	55	blastx.2	G-protein coupled receptor, SREB1 - human	pir JC7287 JC7287	57%	3	368
НОДВF86	859572	99	blastx.2	(AF123880) gag polyprotein [multiple sclerosis associated retrovirus element]	gb AAD48374.1	40% 35%	225 542	437
НОАВР9 0	859622	57	blastx.2	(AF089895) cAMP-dependent protein kinase subunit R2 beta [Oryctolagus cuniculus]	gb AAF00037.1	%26	628	735
HE8UE42	862010	58	blastx.2	(AJ224875) glucosyltransferase [Homo sapiens]	emb CAA12176.1	%66	15	1610
HUSJJS9	862481	59	blastx.2	(AF070661) HSPC005 [Homo sapiens]	gb AAD20967.1	100%	08	316
HSSDM07	863515	09	blastx.2	TcD37 homolog [Homo sapiens]	gb AAF04914.1 U670 85_1	94% 95% 95%	224 647 160	691 1090 225
HBCBW52	866444	61	blastx.2	(AK000385) unnamed protein product [Homo sapiens]	dbj BAA91131.1	%69	1498	1202
HOGAS18	696198		blastx.2	(AK000496) unnamed protein product [Homo sapiens]	dbj BAA91205.1	%6 <i>L</i>	508	380
HHSDL18	869701	63	blastx.2	(AF118082) PRO1902 [Homo sapiens]	gb AAF22026.1 AF1 18094_21	65% 61% 67%	590 429 317	721 536 427

798	628		911	643		397		763			373	283	454	1	044	<u></u>	504		1	103	18	7.67		
1232	173		156	44		104		170	<u>.</u>		23	7	350		m .		4			207	;	69		
100%	100%		 %86	93%		31%		100%			%09	53%	%08		%26		%86			82%		100%		
gb AAD34622.1 AF1	gb AAA35499.1		emb CAA39271.1	emb CAA75163.1		emb CAB70840.1		emb CAA58127.1			dbj BAA23661.1		gb AAC17109.1		dbj BAA91741.1		dbj BAA91018.1			emb CAA59444.1		emb CAA32763.1		
23-H7	Homo sapiens A4 protein [Homo	sapiens]	5'-nucleotidase [Homo	SPIN protein [Homo	sapiens]	(AL137608)	hypothetical protein	neutrophil gelatinase	associated lipocalin	[Homo sapiens]	GCP170 [Homo	sapiens	(AF064604) KE03	protein [Homo sapiens]	(AK001527) unnamed	protein product [Homo	(AK000219) unnamed	protein product [Homo	sapiens	catalase	[Campylobacter jejuni]	propionyl-CoA	carboxylase [Homo	Sapremal
blastx.2	blastx.2		blastx.2	hlastx 2	7 Vigoro	blastx.2		blastx.2			blastx.2		blastx.2		blastx.2		blastx.2			blastx.2		blastx.2		
65	99	}	29	89	ŝ	69		70			71		72		73		74			75		9/		
874506	874608		874787	874801	1,404/0	874930	•	874931			875093		875352		875371		875682			876052		876487		
HLDOG81	HTYOE81		HCRNG90	תסמולעד לא		HWLMV6		HOMANOR			HWLRF06		HODDO41		HCHMQ74		HT3 A 155	2011/2011		HMVBD68		HOCTA74		

HISET05	876686	77	blastx.2	(AJ270993) homeobox	emb[CAB65909.1]	95%	511	849
				protein [Homo sapiens]		%99	801	926
HCRQM22	969928	78	blastx.2	hepatocyte nuclear	gb AAC51129.1	84%	3	401
				factor-3/fork head				
				homolog 11B [Homo 1				
HLHTC92	877310	6/	blastx.2	neuronal PAS2 [Homo	gb[AAB47250.1]	%26	197	418
				sapiens]		%56	142	201
						32%.	482	574
H2CAA49	879484	08	blastx.2	(AF053651) cellular	gb AAC35297.1	100%	954	2108
				apoptosis susceptibility				
				protein [Homo sapiens]				
HETKQ94	880545	81	blastx.2	(AL132954) putative	emb CAB75750.1	35%	53	838
				protein [Arabidopsis		23%	1556	2080
				thaliana]		33%	1155	1298
HOFM023	882466	82	blastx.2	(AF089745) FK506-	gb AAC78853.1	%56	14	499.
				binding protein [Homo		24%	14	499
	•			sapiens]		46%	14	454
						52%	194	499
HNTNP58	882787	83	blastx.2	(AF118082) PRO1902	gb AAF22026.1 AF1	62%	2230	2391
				[Homo sapiens]	18094_21	%09	2382	2480
HOFNY28	888480	84	blastx.2	AF-1 [Homo sapiens]	gb AAA16955.1	93%	18	476
			•			62%	416	658
HE8MQ01	889128	85	blastx.2	(AF118082) PRO1902 [Homo sapiens]	gb AAF22026.1 AF1 18094 21	71%	2699	2490
HE2RG21	891139	98	blastx.2	peptidylprolyl	emb CAA37039.1	100%	59	553
				isomerase [Homo				
	•			sapiens]				•
HHENW77	894855	28	blastx.2	vacuolar proton-	emb CAA75571.1	71%	66	338
				A I Fase submit M9.2				

				[Homo sapiens]				
HOFNU55	897344	88	blastx.2	(AP000616) similar to	dbj BAA85438.1	70%	654	244
				RING-H2 finger				
				protein KHAIa	-	:		
			-	(AFU/8083) [UIYZa safiya]				
TIEOTEAN	6927862	0%	blastx.2	Rabin3 (Rattus	gb AAA67890.1	87%	152	1114
neo Leav	700/60	3		norvegicus		93%	1086	1532
HNTCH73	900546	06	blastx.2	bone morphogenetic protein 2B [Homo	gb AAA51835.1	100%	195	1418
			-	sapiens]			†	900
THIED VS2	002534	9	hlastx.2	IDN4-GGTR14	sp Q9Y6Y5 Q9Y6Y5	95%	0	120
HELD I 33	+55706	1		PROTEIN.		75%	837	926
						20%	801	932
						22%	833	940
						70%	836	955
	272700	5	Llooty 2	(A F200706) TINC-84A	gblAAF15883.1 AF2	47%	1852	2418
HDPIE44	904/03	7,	Diasta.4	[Caenorhahditis	00706 1	39%	170	313
				olongus]	1	25%	1375	1524
				cregame		45%	109	171
LILLION VIO	004783	8	blastx.2	delta-9 desaturase	emb CAA42997.1	71%	6	089
VIACULI		<u> </u>		[Gallus gallus]				01.0
LIDIA V92	906066	76	blastx.2	weakly similar to E.	gb AAA81143.1	36%	1167	2459
COVERIE				nidulans bimA gene		37%	288	1025
				product (SP:P17885)		19%	1776	2474
				Losnorhobditis		22%	1716	2186
			·	[Cacholinacenes		21%	1653	2093
				ciegaio		24%	2274	2540
HOFNG28	908746	95	blastx.2	CD59	sp O62680 CD59_PI G	24%	51	401

	_	380 745	329 427	11 1498		1122 874		53 232	1614 1754		423 364					423 364			309 229	332 249	819 938			064 677	
,	%16	17%	42%	100%		23%		100%	100%		%56	100%	100%	100%	77%	%06	84%	44%	33%	35%	%08	 -	1003	0,70	
	emb CAB43384.1			gb[AAA67038.1]		gb AAF24054.1 AF0	90942 1	dbj BAA88422.1			dbj BAA85438.1										dbj BAA91131.1		10 000000	emo CAA20980.2	
PRECURSOR (MEMBRANE ATTACK COMPLEX 1 (PROTECTIN).	(AL050283)	hypothetical protein	[Homo sapiens]	initiation factor 2	[Homo sapiens]	(AF090942) PRO0657	[Homo sapiens]	(AB029042) ATPase	inhibitor precursor	[Homo sapiens]	(AP000616) similar to	RING-H2 finger	protein RHA1a	(AF078683) [Oryza	sativa	•	·				(AK000385) unnamed	protein product [Homo	Sapiens	(AL031629) similar to	(aka RRM, RBD, or 1 1
٠	blastx.2			blastx.2		blastx.2		blastx.2		-	blastx.2										blastx.2	-		blastx.2	
	96			97		86		66			100										101		,	102	
	911947			913831		914163		915068			915214										917394			91//16	
	HOFOB27			HHBHJ45		HHPDV90		HWMMQ4	7		HBGNY08										HODGH45			HWLKES8	

942		1129	386		492	953	102	808	385	1000	033	,	1/4	245	433	336	781	1293	489	867	607	971	2399
37		113	156		187	687	1	384	7	1	44./	90	77	126	344	307	1281	1364	Ţ	709	329	876	2031
100%		85%	54%		35%	30%	00.20	%86	%66		100%		%09	1007	40%	20%	100%	73%	47%	20%	31%	40%	100%
emb[CAA81066.1]		gb AAC37547.1	emblCAA45262.1		gb AAB01069.1			emb[CAB70771.1]	-		gb AAA58350.1		gb[AAA88036.1]	11 07 700 7 1011	emb[CAA3/04/.1]		ob AAC26082.1		gb[AAC51269.1]				dbj BAA91234.1
dodecenoyl-CoA delta-	isomerase [Homo	cathepsin B [Homo	Sapiens	[Phoca vitulina]	inositol polyphosphate	4-phosphatase [Rattus	norvegicus]	(AT.137496)	hypothetical protein	[Homo sapiens]	replication protein A 14kDa subunit [Homo	sapiens	unknown protein	HOURS Sapiens	ORF4 [Rattus	norvegicus]	(A E034633) GPR 30	(A1.03-033) Of 10.	mitative n150 [Homo	putuity proc process	[crrordes		(AK000530) unnamed protein product [Homo
blastx.2		blastx.2	Flooty 2	Ulasta.4	blastx 2	2		hlacty 2	Tier Carlo		blastx.2		blastx.2		blastx.2		1,10,047, 0	DIASTX.2	blocty 2	Diasta:			blastx.2
103		104	105	3	106	3		107	2		108		109		110			111	113	711			113
919433		919895	220000	607076	2021207	771101		000101	161776		922879		922955		925296		2 (24 ()	925735	200300	158576			926067
HRADL40		HOFMP73		HOFMS02	נותוו/וווו			十	HIMVDQ4	4	HODGI03		HODHD14		HODDC03			HVCAG04		HCKCE02			HHERA91

HAGDL82 928 HWBDS34 930				saniens				
	928396	114	blastx.2	(AK000496) unnamed protein product [Homo sapiens]	dbj BAA91205.1	75% 75%	210	208 257
	930484	115	blastx.2	Similar to S. cerevisiae hypothetical protein L3111 (S59316) [Homo sapiens]	dbj BAA13405.1	%66	57	983
H6ESA39 933	933568	116	blastx.2	BCDNA:LD21720 PROTEIN.	sp Q9XZ25 Q9XZ25	47% 26%	623 14	1150 169
HPCOR06 933	933678	117	blastx.2	ORFII [Homo sapiens]	emb CAA36480.1	73%	310	266 459
						%09 809	266	310
HTFNO06 933	933989	118	blastx.2	(AC009177) unknown protein [Arabidopsis thaliana]	gb AAF27013.1 AC0 09177_3	39%	95	505
HODFY08 935	935520	119	blastx.2	(AB012223) ORF2 [Canis familiaris]	dbj BAA25253.1	45%	79	447
HRACK27 935	935717	120	blastx.2	(AL133045) hypothetical protein [Homo sapiens]	emb CAB61372.1	%26	681	1283
HRACW30 940	940639	121	blastx.2	(AF11387) kappa 1 immunoglobulin light chain [Homo sapiens]	gb AAD29608.1	93%	79	786
HOFOC32 941	941969	122	blastx.2	placental protein Diff33 [Homo sapiens]	emb[CAB09783.1	84%	84	1502
HTTDM15 942	942511	123	blastx	(AL033127) hypothetical protein	emb CAA21869.1	27%	62 759	427 959

	-		Schizosaccharomyces		32%		0101
			pompe	emblCA A43593.11	%59	583	488
944031	124	blastx.2	L1 retroposon, a portion of its ORF2		43%	229	125
			sequence [Rattus		34%	492	0/5
			norvegicus	11 00000	7070	409	2072
946502	125	blastx.2	(AL050294)	emb CAB43393.1	88%	1984	2037
			hypothetical protein		52%	2039	2113
048608	126	blastx.2	ORF2, encodes a	gb AAB60345.1	%09	66	182
)			reverse transcriptase		%10	257	310
			homolog [Homo		64%	169	243
			sapiens	-LIA A A 27565 1	46%	177	365
948708	127	blastx.2	envelope polyprotein	golden strategist	45%	64	174
			[Mus musculus]		85%	7	61
01001	900	blocky 2	(AE038963) RNA	gb AAD19826.1	%66	3	1343
948/19	071	01astA:4	helicase [Homo				
			sapiens]		,000	9	125
949001	129	blastx.2	IDN4-GGTR14	sp Q9Y6Y5 Q9Y6Y5	93%	04	CC1
			PROTEIN.	-ti A A 51851 11	46%	72	539
949574	130	blastx.2	human complement		3/9/	604	732
			Clr [Homo sapiens]		25%	567	740
050057	131	hlastx.2	(AB023584) reduced	dbj BAA88923.1	100%	29	919
70700			expression in cancer				
			[Homo sapiens]		,000	C	536
950475	132	blastx.2	(AF124522) tetraspan NFT-2 [Homo sapiens]	gb AAD17317.1	100%	<u> </u>	000

HCCMDSS 9:	568956	133	blastx.2	catalase	emb CAA59444.1	%46	225	115
1				[Campylobacter jejuni]				
8	859096	135	blastx.2	(AF179370) insulin-	gb AAD52683.1 AF1	%46	424	1551
				like growth factor	79370_1	70%	113	499
				binding protein 5		32%	6	155
				protease [Rattus norvegicus]				
18	961337	136	blastx.2	proline-rich protein	gb AAA53048.1	31%	515	1117
				[Mus musculus]		33%	515	1117
						33%	645	1397
				-		33%	809	1117
				,		33%	1099	1452
						27%	1099	1557
12	961458	137	blastx.2	(AL133051)	emb CAB61378.1	100%	219	1028
				hypothetical protein [Homo sapiens]				
18	962362	138	blastx.2	unknown protein	gb AAA88036.1	%09	1969	1805
				[Homo sapiens]		38%	1820	1680
HBCJN16 96	965190	139	blastx.2	calcyphosine [Homo sapiens]	emb CAA66609.1	%66	38	346
8	965372	140	blastx.2	(AF083217) WD repeat	gb AAD45865.1 AF0	%/6	32	1660
				protein WDR3 [Homo	83217_1	27%	473	940
				sapiens]		53%	2270	2308
				•		34%	2226	2312
						%09	2269	2313
		_				%09	2273	2317
•		_	_			%09	2272	2316
						46%	5269	2307
•					-	36%	2226	2315
·						53%	2270	2314

(AK000069) unnamed protein product [Homo sapiens]
similar to
D.melanogaster
peroxidasin(U11052) [Homo sapiens]
(AB017644) ubiquitin-
conjugating enzyme E2 [Homo sapiens]
preceruloplasmin (EC
1.16.3.1) [Homo
(AK000200) unnamed
protein product [Homo
F20D12.3 gene product
Caenorhabditis
hPMSR6 [Homo
SELENOPROTEIN
PRECURSOR.
(AB012955) KIP2
Homo sapiens
hepatocyte growth
돰

				factor-like protein		%05	100	579
			·	[Homo sapiens]		44%	100	579
HWLHW8 6	975771	153	blastx.2	epithelial glycoprotein (EGP) precursor Homo saniens	gb AAA35723.1	100%	245	1186
HOFNM53	976051	154	blastx.2	(AL110276) hypothetical protein Homo saniens	emb CAB53711.1	37%	498	1094
HDPSE86	976207	155	blastx	(AF086713) rasGAP- activating-like protein Homo sapiens	gb AAD09006.1	83% 91% 96%	757	1311 420 774
						30%	361	511 450 42
HHFOE18	976216	156	blastx.2	(AL117664) hypothetical protein [Homo sapiens]	emb CAB56034.1	52% 35%	376 14	1275
HHFNH27	896926	157	blastx.2	Nascent polypeptide associated complex alpha subunit [Homo sapiens]	emb CAA56869.1	98% 100%	33	589 26
HMMBZ8 1	977264	158	blastx.2	alternatively spliced product using exon 13A [Homo sapiens]	gb AAB49034.1	52%	545	369
HSLGF32	977704	159	blastx.2	(AL133063) hypothetical protein [Homo sapiens]	emb CAB61387.1	22%	2	250
норғи73	978812	161	blastx.2	myeloid ecotropic viral integration site-1b	gb AAA85509.1	94%	371	652

000	183	1603		626	692	165	204	482	673	620	1659	1262	2071						•	2157	4	88	822
,	365 130	53		136	236	1	4	613	783	299	937	5	301	1207	1684	1702	1609	1711	1819	1867	173	~	727
	38%	% 5 66	200	100%	1000%	40%	45%	72%	70%	%89	100%	,	44%	36%	78%	79%	31%	22%	35%	23%	%69	71%	20%
	gb AAD40191.1	gb AAB58817.1	dbj BAA91821.1	gb AAB88182.1	11000001	emb[CAAoouvu.1]		dbilBAA91131.1			emb CAA62350.1		gb AAD55431.1 AF1	81645 1	1						amblCAR05030.1		
Mins musculus]	356) putative licase [Homo	廿	(AK001665) unnamed protein product [Homo	(AF035299) similar to GAP binding protein	p62do [Homo sapiens]	kinase A anchor protein	ferrordus ornou)	(AVOO0285) unnamed	(Anchorage) unimized	protein product [1101119: 1	alphaII spectrin [Rattus	norvegicus	(AF181645)	Bonna GH12144	DODING: CITED !	Drosopmia	melanogaster]				11.4.4.4.5.5.5	predicted using	to E.coli guanosine-3',
	blastx.2	blastx.2		blastx.2		blastx.2			DIaSTX.2		blocky 2	Olasta:	block 2	Ulasta.2						•	١	blastx.2	
	162	163	164	165		166			167		150	001	160	107								170	
	866826	979346	979468	979547		999626			980139		0,000	697086	000.00	670186						·		981108	
	HNBUA49	HVVD[173	+	HOCPY88		HOGDC64			HSIEA14			HSPSY43		HSXBHZ4								HOFAE61	

				1 [Caenorhabditis				
HCFOF82	981272	171	blastx.2	(AL133558)	emb CAB63713.1	%86	2	901
				hypothetical protein				
HOCMT79	981309	172	blastx.2	similar to yeast Sec6p,	gb AAA85505.1	94%	481	1413
				Swiss-Prot Accession	-	%96	2	481
				Number P32844; 1 1				
				norvegicus				
HKAE103	981319	173	blastx.2	Wiskott-Aldrich	gb AAC52556.1	35%	392	9
				Syndrome Protein [Mus		39%	233	9
				musculus]		40%	407	300
-	-					82%	306	253
						34%	288	211
HOCP031	981593	174	blastx.2	(AF015037)	gb AAB99905.1	%6 <i>L</i>	381	812
				endooligopeptidase A				
				related protein; EOPA				
				related protein				
			•	[Oryctolagus				
		-		cuniculus]				
HAOTG88	981606	175	blastx.2	No definition line	gb AAC43132.1	100%	217	2
				found [Escherichia				
				con				
HVCAH21	981768	176	blastx.2	(AF044956)	gb AAD42057.1 AF0	100%	80	919
				NADH:ubiquinone	44956_1			
				oxidoreductase B22				
				subunit [Homo sapiens]				
HFPCK56	981812	121	blastx.2	(AF099664) Cdc42	gb AAD16299.1	%88	3	125
				effector protein 4				_
				[Homo sapiens]				•

482 28	250	247	245	285	284	283	280	282	281	277	279	278	275	274	276	271	273	272	270	268	569	265	566	267	262	263	264
24	321	318	316	350	349	348	345	347	346	342	344	343	340	339	341	336	338	337	335	333	334	330	331	332	327	328	329
87%	%56	91%	95%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%	100%
emb CAB75427.1	dbilBAA85438.11																										•
(AJ271784) chromokinesin [Homo	sapiens	P.D. C. L. Garage	KUNG-112 111861	VAEOT8682) [Oeuza	(AFU/8082) [O1)24	[avii va														-							
blastx.2	1.10.41. 0	DIASUX.2						*																			
178		4/1								_																	
981859	00000	981862																									
HAOSJ58		HOPJT48																									

259 260 242	241	239	217	1091	156	578	635	942	1137	552						949	157	737	790	705	3170	1451	1451
324 325 313	318	316	318	54	593	763	763	1088	628	121				-		503	∞	63	869	100	2124	1167	1167
100% 100% 91%	%88	84%	64%	%66	36%	33%	37%	29%	48%	46%						%66	%06	%96	74%	100%	32%	38%	38%
				dbj BAA91241.1	emb CAA91932.1		_		emb CAA19455.1							dbj BAA91222.1		emblCAA12271 11		gb AAA36158.1	gb AAB66420.1		
				(AK000541) unnamed protein product [Homo saniens]	similar to cuticle	collagen	[Caenorhabditis	elegans]	(AL023828) cDNA	EST yk289g5.5 comes from this gene; cDNA	EST 1 1 yk653f1.5	comes from this gene;	cDNA EST	EMBL:C07875 comes	from th	(AK000516) unnamed	protein product [Homo	(AJ224979) MTWR1	[Homo sapiens]	L6 [Homo sapiens]	unknown [murine	herpesvirus 68]	
	-	•		blastx.2	blastx.2				blastx.2							blastx.2		hlastx.2		blastx.2	blastx.2	•	,
				180	182				184					_		185		186		187	188		
				981914	982032				982197							982465		982618		982764	800£86		
				HCFAV61	HOVJY54				HE8MM52							HJBCC19		HODAA93		HSPSI74	HCEHZ42		

	Γ			_	г—	·	$\overline{}$		_		т		1				_	_		_
451	626 244	393	369	336	10	069	547	Ì	1202	391	511		1471	1387			591	363	089	1312
185	393 110	238	244	247	096	. 169	37	5	204	7	134		14	14	-		-	103	651	,
.100%	42% 46%	46%	78%	35%	28%	100%	1,000	8,001	7080	97%	31%		54%	53%			100%	48%	70%	100%
emb CAA37064.1	emb CAB45690.1				emb CAA84230.1	emb CAA53619.1	2k A C26220 11	golwyddiad	AbilB A A 23673 11		dbj BAA83500.1		gb AAD09819.1	-	•		emb CAA65246,1	emb CAA39297.1		gb AAA36584.1
t-complex polypeptide 1 (AA 1-556) [Homo sapiens]	(AJ243177) Xenopus RPA interacting protein	alpha [Xenopus laevis]			extensin-like protein [Zea mays]	neuromedin U [Homo	(A E097125) E1EO #ma	ATPase subunit d Homo saniens	GTRP-AIT (Homo	sapiens	(AB031292)	proteolipid protein 2	(AF111423)	chromosome	condensation protein	XCAP-G [Xenopus	erm [Homo sapiens]	put. ORF [Homo	sapiens]	replication protein A.
blastx.2	blastx.2				blastx.2	blastx.2	Floate 2	7.07620	hlacty 2		blastx.2		blastx.2				blastx.2	blastx.2		blastx.2
191	192				193	194	105	3	106	.	197		198				199	200		201
984168	985043				985280	985323	085401	10+000	985580		820986		986158			•	986165	986328		986744
HDABW5 0	НАОВНП				HMVAW4	HAGDF03	HODVIDO	Carrier I	HA ICA 11	5	HWAHA1	-	HSAMI43				HNFJH73	HNTCH03		HSUAA20

1347	1054	+ col		1092 743	0	01/				820	00,	23	1289	476	1238	857	1061	812	428	1278	
1315	\perp	5	İ	256	\perp	8/7				368	-0,	0 m	63	81	699	492	801	516	195	1228	
100%	,800,	100%		76%	2 3	46%	•		 	100%	1,000	%88 88	7070	47%	41%	44%	36%	33%	37%	61%	•
		emb CAB37641.1		emb CAB57330.1		gb AAC61698.1		-		emb CAA69255.1		gb AAD27835.1 AF1 21862_1		emb CAA/5103.1							
70-kDa subunit [Homo		663) 17.6 (Major	Epididymis-specific	<u> </u>	hypothetical protein [Homo sapiens]		spningosine kinase [Mus musculus]			helix-loop-helix protein	[Homo sapiens]	(AF121862) sorting nexin 13 [Homo	sapiens]	SPIN protein [Homo	sapiens]					·	
		blastx.2		blastx.2		blastx.2				hlastx.2		blastx.2		blastx.2		_					
		202		203		204				205	204	206		207							
		292986		987018		987071				007117	70/117	987165		987262							
		HSPAD08		HFKBA32		HHFLU49					HOENAIO	HTFOW71		HTTAG03						- ,	

blastx.2 elongation factor 2 emb CAA35829.1 [Homo sapiens]
blastx.2 (AF134726) NG23 gb AAD21821.1
(AF134/26) NG23 [Homo sapiens]
blastx.2 3-oxoacyl-[acyl-carrier- dbj BAA35901.1 protein] reductase (EC 1.1.1.100). [Escherichia coli]
blastx.2 nucleoporin-like emb CAA61667.1 protein [Homo sapiens]
blastx.2 TBX2 [Homo sapiens] gb AAA73861.1
blastx.2 (AL110239) emb CAB53690.1
hypothetical protein [Homo sapiens]
blastx.2 (AB006572) RPB5 dbj BAA34781.1
meidating protein [Homo sapiens]
blastx.2 cDNA EST yk575f9.3 emb CAA94859.1
comes from this gene
[Caenorhabditis
elegans
blastx.2 envelope protein gb AAA88027.1
[Homo sapiens]
blastx.2 RNA polymerase II gb AAA75522.1
transcribnon factor 5111

blastx.2 blastx.2 blastx.2	high mobility group protein 2a [Homo sapiens] MHC Class I region proline rich protein [Homo sapiens] binding protein [Homo sapiens]	emb CAA71143.1 gb AAB40147.1 emb CAA00862.1	%86 %66	1	366
blastx.2 blastx.2	lass I region rich protein sapiens] protein [Homo	gb AAB40147.1 emb CAA00862.1	%86		
blastx.2	binding protein [Homo sapiens]	emb CAA00862.1		429	728
			99% 52% 56%	153 308 804	587 907 956
			28%	67	222
			31% 45%	637	789
blastx.2	(AJ252060) TRABID protein [Homo sapiens]	emb CAB64449.1	%16	1	2127
blastx.2	Berg36 [Homo sapiens]	emb CAA67781.1	100%	49	789
blastx.2	(AK000385) unnamed protein product [Homo	dbj BAA91131.1	60% 77% 57%	471 301 1148	292 167 1107
blastx.2	(AB007619) EBAG9 [Homo sapiens]	dbj BAA22572.1	100%	535	1173
225 blastx.2	Leu2 [Homo sapiens]	emb CAA75516.1	93% 100% 72%	312 551 260	410 607 346
226 blastx.2	(AB002533) Qip1 [Homo sapiens]	dbj BAA19546.1	92% 24% 47%	120 279 1 5	1178 974 216 121
1 1 1	blastx.2 blastx.2 blastx.2		(AB007619) EBAG9 [Homo sapiens] Leu2 [Homo sapiens] (AB002533) Qip1 [Homo sapiens]	(AB007619) EBAG9 dbj BAA22572.1 [Homo sapiens] emb CAA75516.1 (AB002533) Qip1 dbj BAA19546.1 [Homo sapiens] dbj BAA19546.1	(AB007619) EBAG9 dbj BAA22572.1 100% [Homo sapiens] emb CAA75516.1 93% Leu2 [Homo sapiens] dbj BAA19546.1 92% [Homo sapiens] 47% [Homo sapiens] 35%

247	282	099	618	652	813	790		1155	528	181	542	706	88	2745	330	330	327	330	327.	330	330	330	330	339	327	327
11	250	100	361	2	622	7		616	187	116	282	123	56	43	40	4	43	49	43	46	46	46	46	49	49	43
94%	100%	100%	%26	%86	%68	100%		100%	93%	100%	100%	78%	21%	%86	43%	40%	45%	46%	44%	47%	48%	45%	46%	43%	44%	41%
emb CAA43200.1		dbj BAA01374.2	gb AAF36161.1 AF1 51075 1	gb AAC79695.1		dbj BAA91941.1		gb AAF32373.1 AF2	22742 1		gb AAF32373.1 AF2	22742 1	l	emb CAA36267.1												
transcription factor ILF	-	p67 myc protein	(AF151075) HSPC241 [Homo sapiens]	(AF067817) VAV-3	protein [Homo sapiens]	(AK001851) unnamed	sapiens]	(AF222742) synaptic	glycoprotein SC2	[Homo sapiens]	(AF222742) synaptic	glycoprotein SC2	[Homo sapiens]	collagen type VI. alpha	3 chain [Homo sapiens]	7					•					
blastx.2		blastx.2	blastx.2	blastx.2		blastx.2		blastx.2			blastx.2			blastx.2												
1227		228	229	230		231		232			233			234	}											
209686		989952	090066	990146		990184		990254			990255			990435												
HSPSH36		HAAAA25	HTEMJ16	HWLAB90		HOSED43		HNODE 50			HLWBV17	_		ууно ОН			•									

330	327	327	333	327	096	1587	647	1031	-	3258	202	499				2098			226	63	324	72	72	75	72	75	72
85	46	49	46	49	391	1117	531	75		10	131	107				1436			65	43	241	4	13	-	-	13	4
48%	41%	41%	41%	43%	79%	79%	45%	%66		%26	%99	37%				%60	2		47%	71%	100%	39%	45%	32%	36%	33%	21%
								gb AAA36407.1	•	dhilBAA13432.1		gb AAA83581.1				110 4 10 500 40 11	emo CAB33248.1		1.1 A A COACA 11	golacasostos.1	gb[AAD09746.1]						
		-						pyrroline-5-carboxylate	reductase [Homo	Sapiens	normedicus	coded for by C. elegans	cDNA yk38h3.5; coded	for by C. elegans 11	[Caenorhabditis	elegans	(AL122073)	hypothetical protein	Homo sapicina	ORF 3 [Homo sapiens]	(AE065301) ZIS1	Tiomo canienal	Littomo sapiemaj				
-				•				blastx.2			Diastx.2	blachy 2					blastx.2			blastx.2	110041	DISSIX.2					
								235			730	227	<u>`</u>				238			239	9,0	740					
		,					***	990546			609066	000011	110066				990751			692066		990913					
								HOGDC67			HCDBO02		HODGINAZ				HPDRP30			HBXFN09		HDTB075					

1-0	000	0 m 0	Ø.	24	0	3	ω Ø.		0	~
902 181 470	1550 1710 1929	492 873 940	449	7	059	333	903 1079	811	540	407
192 32 171	1104 1552 1747	238 607 803	. 192	263	210	40	598 933	8	157	111
47% 52% 38%	42% 56% 27%	100% 92% 69%	100%	76%	100%	%96	100% 79%	%19	%66	48%
gb AAD23834.1 AF1 23653_1	gb AAF23786.1 AF2 13822_1	gb AAC51733.1	dbj BAA91655.1	dbj BAA91131.1	gb AAC51768.1	dbj BAA07598.1	dbj BAA91189.1	gb AAF09133.1	gb AAF36161.1 AF1 51075_1	emb CAB04720.1
(AF123653) FEZ1 [Homo sapiens]	(AF213822) hypothetical protein [Zymomonas mobilis]	(AF007872) torsinB [Homo sapiens]	(AK001371) unnamed protein product [Homo sapiens]	(AK000385) unnamed protein product [Homo sapiens]	microsomal glutathione S-transferase 2 [Homosapiens]	TSC-22 [Homo sapiens]	(AK000474) unnamed protein product [Homo sapiens]	(AF176524) F-box protein FBL10 [Mus musculus]	(AF151075) HSPC241 [Homo sapiens]	CDNA EST
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
241	242	243	244	245	246	247	248	250	251	252
991014	991048	991268	991516	991654	991761	992678	992780	992973	082866	993403
HELGN26	HODCUIS	новыя1	HLWAF02	HRKPA16	нерох94	HEAAY09	HPDRH78	HODFOS7	HOCPZ44	HPAMU38

91	372	205 205	134	613	382	671	695	;	1531		295	155	558	536	153	528	499	404	377
20	277	124	m	545	239	624	٣	,	1073		. 155	122	421	147	-	412	413	291	273
37%	78%	64% 65%	48%	%69	%95	20%	430%	8/C+	%28		%50	2000	43%	730%	7070	%95	51%	34%	28%
	gb AAD33288.1 AF1 26163 1	gb[AAC51261.1]			dbilBAA92096.1		11000001011	emb CAA32297.11	dbj BAA92074.1		114 A TOOPOOT 11 A E 1	gb AAU2//2/.1 AF1	32952_1	11 NSC10 A ACT: 16	doj bAA91334.1				-
yk269g12.5 comes from this gene; cDNA EST EMBL:D27364 1 1 comes from this gene; cDNA EST EMBL:D36272 comes	(AF126163) HHLA3	putative p150 [Homo	sapiens		(AK002129) unnamed	protein product [Homo	sapiens	putative [Rattus	(AK002081) unnamed	protein product [Homo	sapiens	(AF132952) CGI-18	protein [Homo sapiens]		(AK000741) unnamed	protein product [Homo	sapiens]		
	blastx.2	blastx.2			1100tr 2	Olasia.4		blastx.2	Linety 2	Ulasta: 4		blastx.2			blastx.2				
	253	254			220	667		256	250	/67		258			259				
	993602	993754				993806		993918	10000	993931		994134			994234				
	HNGGK47	HODGN51			-+	HODCT60		HAQBV81		HDTGF49		CHUGRNES	10001		HSKGR42				

			<u> </u>					
468 437	362	1134	410	531	513	339	1157 270	1311
316 291	3	1	231	169	85	43	273 124	46
28% 32%	%26	100%	100% 81%	100%	100%	%4%	98%	%96
	emb CAB55923.1	emb CAA33389.1	emb CAA38702.1	gb AAC50885.1	dbj BAA03400.1	emb CAB37991.1	emb CAA72364.1	gb AAB51383.1
	(AL117435) hypothetical protein [Homo sapiens]	creatine kinase B	Cks1 protein homologue [Homo sapiens]	NifU-like protein [Homo sapiens]	ribosomal protein [Homo sapiens]	(AL031432) dJ465N24.1 (PUTATIVE novel protein similar to predicted yeast and worm proteins) [Homo sapiens]	phosphate cyclase [Homo sapiens]	lin-10 protein homolog [Rattus norvegicus]
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	260	261	262	263	265	266	267	268
	994356	994536	994596	994664	994776	994874	994954	994993
	ноево85	HOPJG01	HKBAK06	HKGCN61	HFOYI37	HOFNL18	HOFNT57	HCBMT45

	DIASTX.2	(ALUSIOOO) WOTENLA)	emolectory/+1::1	2001	?	
		(eukaryotic translation initiation factor 2,				
		subunit 2 (beta, 38kD))	-			
		[Homo sapiens]	110117741011	7000	×	520
270	blastx.2	(AL050348) dJ447F3.2	emb[CAB66118.1]	9470	o	2
		(ubiquitin-conjugating				
		enzyme 52 m10) [Homo saniens]			·	
271	blastx.2	alpha 4 protein [Homo	emb CAA70119.1	%66	73	981
		sapiens]		2000	27	929
272	blastx.2	proteasome subunit	dbj BAA05647.1	99%	,	900
273	blastx.2	SRp30c [Homo	gb AAA93069.1	100%	100	762
		sapiens		740/	22	140
274	blastx.2	NS GENE PRODUCT LONG ISOFORM,	sp G1488414 G14884 14	63%	5 2	34
		N8L PROTEIN=D52 1				•
١		HN4A1	JE:15 4 401400 1	%0%	66	506
275	blastx.2 	(AK000857) unnamed protein product [Homo	aojibAA31400.1	100%	419	637
		sapiens]		/070	90	1307
276	blastx.2	NF45 protein [Homo	gb AAA20993.1	70%	1175	1297
ı	1.1	DAIA nolymerase II	gb[AAA91459.1]	100%	219	419
117	DIASTX.2	subunit [Homo sapiens]	- 1			S
278	blastx.2	(AB030654) AP-4	dbj BAA82970.1	100%	4	777
	_	clathrin adaptor-related				
		complex sigma4	. •			
- 1		Subunit Indued Sapicus				

222	456	228	352	759 244	735	1174	957 298	1360	380
770	1	1	218	244	301	1037	274 32	1226 502	162
%16	%96	%86	%16	100%	34% 44%	100%	97% 97%	%L6	%08
gb AAA03583.1	gb AAA36585.1	dbj BAA91517.1	gb AAD34147.1 AF1 52097 1	emb CAA99731.1	gb AAC47967.1	gb AAF19255.1 AC0 04858_3	emb CAB55313.1	dbj BAA09762.1	gb AAB42266.1
L-lactate permease Escherichia colil	rac protein kinase-beta	(AK001138) unnamed protein product [Homo saniens]	(AF152097) CGI-05 protein [Homo sapiens]	nuclear protein SA-1 Homo sapiens	C13F10.7 gene product [Caenorhabditis elegans]	(AC004858) U1 small ribonucleoprotein 1SNRP homolog; match to PID:g4050087 Homo sapiens]	(AJ132948) rfg7 protein [Homo sapiens]	product is related to clathrin-associated protein. [Homo sapiens]	weak similarity to Arabidopsis thaliana ubiquitin-like protein 8 [Caenorhabditis
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
279	280	281	282	283	284	285	286	287	288
996874	606966	997127	997165	997595	997862	997873	650866	998517	998533
нненк49	нснсғ36	HAHCK58	HETIJ06	HAPOE30	HRGDC33	HMTMB9 1	HFAAD07	HE8TG67.	HACNC39

	100000	289	blastx.2		dbjjbAA82000.1	2001)	
,				type2 [Homo sapiens]	omblC 4 B 56776 11	100%	47	664
HCGMA67	998905	290	blastx.2	\dagger	11 4 DA 505 11 A CO	%86	2	466
HSKHK19	896866	291	blastx.2	lar to	gp AAD43623.1 AC0	3%%	995	721
				HUB1; similar to	7-04930	54%	1095	1160
-				BAA24380 (PID:g2789430)		70%	558	587
				[Homo sapiens]				200
HAGGR 59	999124	292	blastx.2	histone H1-I [Volvox	gb AAA74723.1	36%	270	986
				carteri]		,000	301	1804
HOPKS83	999148	293	blastx.2	(AF026124)	gb[AAC73069.1]	93%	C74	1071
				schwannoma-		-		
				associated protein [Mus				
				musculus]	spi015352 015352	100%	691	1368
HE8CY70	999157	294	blastx.2	AB A GATENT		95%	32	691
٠				(FICEGIVIENT):		%99	m	128
						48%	635	706
		,	0.11	(A 1050562) tetraspanin	emb[CAB65594.1]	%88	34	546
HPCT153	999243	295	DIASTX.2	(AJ2JUJ02) teuaspaimi		100%	474	779
	_	900	Llocty 2	(AF017790)	gb AAB80726.1	%56	152	1702
HOPKN50	999313	720	Ulasta.4	retinoblastoma-	-			
				associated protein HEC		-		
	-			[Homo sapiens]		, 000	(116
HAPAI17	999778	297	blastx.2	(AK001832) unnamed protein product [Homo	dbj BAA91931.1 	%/6	7	CII
				sapiens	1, 00,000	7007	96	1903
HHAUV59	808666	298	blastx.2	p116Rip [Mus	gplace	%29	547	801
		•	•	[snmosnut		21%	1454	1669

HDPUH64 100033	1	1.7.7	X	(AK001770) unnamed	dhi BAA91897 1	36%	663	992
100				protein product [Homo		27%	999	1379
100				sapiens		792	4	645
9			•	1	•	39%	481	621
	1000339	300	blastx.2	CRAG protein	emb CAA76938.1	45%	∞	869
				Drosophila	-	%99	525	692
				melanogaster	-	65%	475	534
100	1000424	301	blastx.2	(AL117183) conserved	emb CAB54870.1	25%	55	720
				hypothetical protein		36%	755	994
				[Schizosaccharomyces				
				pombe]				
100	1000582	302	blastx.2	(AJ387747) sialin [Homo sapiens]	emb CAB62540.1	100%	3	854
100066	6990	303	blastx.2	(AF176555) A-kinase	gb AAF07045.1 AF1	94%	9/	627
				anchoring protein 220	76555_1			
				[Homo sapiens]				
100	1000875	304	blastx.2	(AK000462) unnamed	dbj BAA91181.1	41%	296	556
				protein product [Homo				·
				sapiens]				
100	1001066	305	blastx.2	(AF047002)	gb AAD09608.1	%86	7	999
				transcriptional				
				coactivator ALY		,	_	
				[Homo sapiens]				
100133	1333	908	blastx.2	(AK000541) unnamed	dbj BAA91241.1	48%	40	237
				protein product [Homo				
				sapiens	-			
100	1001695	208	blastx.2	17-kDa protein [Homo	gb AAA36038.1	%66	127	621
				sapiens]				
100	1001901	308	blastx.2	pol [porcine	emb CAA76582.1	41%	398	721
				endogenous retrovirus]		33%	25	423

807	701	652	639	1041	508	356	342	689	826	106
373	105	23	13	64	353	3	<i>L9</i>	57	134	2
100%	100%	73%	100%	%86	%86	%66	100%	32%	%96	85%
gb AAA89070.1	emb CAA48137.1	dbj BAA91724.1	dbj BAA91787.1	emb CAB58438.1	gb AAA60116.1	emb CAA93701.1	emb CAA31007.1	dbj BAA82444.1	emb[CAA40940.1	sp Q9Y6Y5 Q9Y6Y5
S19 ribosomal protein	proliferation associated gene (pag) [Homo	(AK001496) unnamed protein product [Homo saviens]	(AK001610) unnamed protein product [Homo sapiens]	immunoglobulin gamma-2 heavy chain Homo sapiens]	lysyl hydroxylase [Homo sapiens]	Huntington's Disease (HD) gene [Homo	snRNP E protein (AA 1-92) [Homo sapiens]	(AB022660) SET-binding protein (SEB)	immunoglobulin lambda light chain [Homo sapiens]	IDN4-GGTR14 PROTEIN
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	.blastx.2
309	310	311	312	313	314	315	316	317	318	319
1001954	1002071	1002096	1002328	1002459	1002468	1002479	1002492	1002527	1002545	1002546
HVVCB28	HCOOS01	HDACA35	HOVDG59	HTJAD78	нсрск69	HPTTW90	HSUBG36	HODFU72	HCNSF57	HODJU13

1556	755	823	923	1408	361	681	176	281	659	831	925	768
207	174	407 209	153	98	86	244	6 0, 6	m	591	178	323	28
%66	100%	91% 53%	100%	100%	100%	100%	%85	42%	41%	100%	100%	100%
dbj BAA11748.1	gb AAC83802.1	emb CAB69299.1	gb AAA35822.1	emb CAA45089.1	gb AAA60280.1	gb AAA60583.1	gb AAB24264.2			gb AAA59505.1	dbj BAA05646.1	gb AAD15546.1
nuclear protein, NP220 [Homo sapiens]	(AF097441) phenylalanine-tRNA synthetase [Homo sapiens]	unnamed protein product [unidentified]	folate-binding protein precursor [Homo sapiens]	homologue to elongation factor 1- gamma from A.salina [Homo sapiens]	ribosomal protein L37a [Homo sapiens]	RPS16 [Homo sapiens]	HKR-T1 [Homo	sapiens]		clathrin light-chain A [Homo sapiens]	proteasome subunit HsC7-I [Homo sapiens]	(AC004983) similar to PID:g3877944 [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2			blastx.2	blastx.2	blastx.2
320	321	322	323	324	325	326	327			328	330	332
1002551	1002563	1002565	1002591	1002607	1002610	1002729	1002807			1002811	1003155	1003224
HODIL25	ночнѕѕб	HPMJQ18	HLHC146	HAOSD18	HACNG47	HOPJX95	HSCLR05			HVVAO74	HVVBK18	HSCLM55

protein 73% 73% 100% 28% 100% 28%			•	t		emblCAB63754.1	%66	881	1477
Hono sapiens Hypothetical process Hono sapiens Hono sapiens 173% 548 518	\vdash			_			%66	504	881
Homo sapiens Homo sapiens Homo sapiens 100% 373 373 500	_				hypothetical process		73%	40	510
1004480 334 blastx.2 (AF026692) ftpHE gb AAC04617.1 93% 878 108 108 1004480 334 blastx.2 (AF026692) ftpHE gb AAC04617.1 93% 135 135 145 13685 141% 1004619 336 blastx.2 (AF113685) PRO0974 gb AAF29584.1 AF1 56% 171 2 265 33 141 66 20 2 20 2 20 2 20 2 2	-				[Homo sapiens]		100%	373	207
1004480 334 blastx.2 (AF026692) ftpHE gb AAC04617.1 93% 132 132 45 45 45 45 45 45 45 4							28%	878	1084
1004480 334 blastx.2 (AF026692) ftpHE gb AAC04617.1 93% 18 51	•						79%	1121	1327
1004480 334 blastx.2 (AF026692) ftpHE gb AAC04617.1 93% 18 51							35%	343	459
1004639 334 blastx.2 (APC200223) appears 1004583 335 blastx.2 (AB012223) ORF2 dbj BAA25253.1 41% 362 76 1004583 335 blastx.2 (AF113685) PRO0974 gb AAF29584.1 AF1 56% 171 2 200 21 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200 22 200			一	,	Hung (COSSCOULAN	gb[AAC04617.1]	93%	18	515
1004583 335 blastx.2 (AB012223) ORF2 dbj BAA25253.1 51% 35 135 135 135 136 13		1004480		blastx.2	(AF020032) APACE	- 0	/01/	26.2	760
1004585 355 1004619 336 blastx.2 capitalistis dip BAA21615.1 69% 171 2 1804619 336 blastx.2 capitalistis dip BAA21615.1 69% 171 2 1804625 337 blastx.2 capitalistis dip BAA21615.1 69% 10 1 1004621 338 blastx.2 capitalistis dip BAA21615.1 69% 10 1 1004621 338 blastx.2 capitalistis capitalis capi	-	2007	十	hlactx 2	(AB012223) ORF2	dbj BAA25253.1	41% %1%	302	143
1004619 336 blastx.2 (AF113685) PRO0974 gb AAF29584.1 AF1 56% 171 2 200		1004583			Canis familiaris]		83%	135	191
1004619 336 blastx.2 (AF113685) PRO0974 gb AAF29584.1 AF1 56% 171 2 38% 277 4 4 4 4 4 4 4 6 6 6					•		51%	265	360
1004619 336 blastx.2 (AF113685) PRO0974 gb AAF29584.1 AF1 56% 171 2 38% 277 4 4 4 4 4 4 4 4 4							70%	200	259
1004619 336 blastx.2 (AF113685) PRO09/4 golyara 273 control of the property of the property of the property of the protein product [Homo sapiens] 13685 1 100% 141 65					7 6000 110	11 A A E20584 11AF1	%95	171	293
Homo sapients Sapients Sapients Homo s	4741UOD	1004619	336	blastx.2	(AF113685) PRO09/4	golder 22307.11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	38%	277	453
1004625 337 blastx.2 putative [Homo solitor 1004627 338 blastx.2 AB005878 BYJ15 dbj BAA21615.1 69% 10 1	S/ANGON				Homo sapiens	mb A C37567.1	100%	141	683
1004627 338 blastx.2 (AB005878) BYJ15 dbj BAA21615.1 69% 10 10	HODIVED	1004625	337	blastx.2	putative [Homo	Sologia Coversion			1
1004627 338 blastx.2 (AB0028/8) B 1 1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	201 CTOU				sapiens	ANIBA A21615.11	%69	10	135
1004631 339 blastx.2 DNA primase (p58 emb CAA52378.1 90% 1 1004632 340 blastx.2 (AF027728) kinesin- Eated protein (AF027728) kinesin- Eated protein (AF027728) kinesin- Eated protein 24% 524 1 24% 524 1 Eated protein Eated	HODEF10	1004627	338	blastx.2	(AB005878) B 1313	and the second s		1	780
1004631 339 blastx.2 blastx.2 subunit) [Homo sapiens] 25% 518 1 1004632 340 blastx.2 (AF027728) kinesin- 24% 524 1	 				DATA mimase (n58	emb CAA52378.1	06		707
1004632 340 blastx.2 (AF027728) kinesin- gb AAC60300.1 25% 518 1	HODFB03	1004631	339	Diastx.2	subunit) (Homo				
1004632 340 blastx.2 (AF027728) kinesin- gb AAC60300.1 25% 135 135 136 135 136					saniens		1,050	510	1165
1004632 340 Diasta. related protein 24% 524 1			(,	Lingto 2	(AF027728) kinesin-	gb AAC60300.1	72%	135	533
[Xenopus laevis] 23% 524 1 1004633 341 blastx.2 protein product [Homo sapiens] sapiens]	HPMLN08	1004632	340	Jan Viene	related protein		26%	524	1051
1004633 341 blastx.2 (AK001410) unnamed dbj BAA91675.1 87% 22 protein product [Homo sapiens]					[Xenopus laevis]		23%	524	1060
1004633 341 blastx.2 (AK001410) unnamed db] BAA3107231 protein product [Homo sapiens]						11 777 10 A A CIL 11	87%	22	369
sapiens	НОДЕН08	+	341	blastx.2	(AK001410) unnamed protein product [Homo				
					sapiens				

758	142	208 2175 2150 2053	236	1238 1245	1292	400	963	453	453 444	408	453	453	276
165	405	68 1993 2055 2030	3	435 1204	153	185	13	-1 -4	91 4	4	4		4
%86	52%	97% 52% 62% 87%	100%	97% 66%	100%	%08	%68 %68	%65 29%	53%	28%	48%	43%	65%
gb AAC39906.1	dbj BAA91205.1	emb CAB52687.1	dbj BAA91965.1	emb CAA57684.1	gb AAA52471.1	gb AAA52360.1	pir S26650 S26650						
(AF069736) PCAF associated factor 65 beta [Homo sapiens]	(AK000496) unnamed protein product [Homo sapiens]	(AJ010089) GANP protein [Homo sapiens]	(AK001892) unnamed protein product [Homo sapiens]	gpStaf50 [Homo sapiens]	c-fos protein [Homo sapiens]	Ral guanine nucleotide dissociation stimulator [Homo sapiens]	DNA-binding protein 5 - human						
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2						
342	343	344	345	346	347	348	349						
1004635	1004643	1004648	1004660	1004887	1004950	1005153	1005236						
HEGBF25	норъ	норесз8	HODIQ24	HVVDD56	HTHDV01	HVVDJ95	HODIY29						

444	441	1013	1015	688	889	111	749	737	740	/40	205		403		2081	381 448	509			1037 603
1	25	. 309	2	107	155	58	069	717	720	717.1	89	1	7		411	377	435		·	573 136
41%	45%	71%	100%	97%	33%	%88	70%	100%	100%	87%	%91		% 86		%66	98% 54%	100%			80%
		gb AAB50568.1	gb AAA75595.2	gb AAD00897.1							emb CAB09660.1		gb AAD40846.1 AF0 72441_1		dbj BAA91829.1		gb AAB49678.1	,		gb[AAC50697.1
		uridine kinase [Mus	tax1-binding protein TXBP151 [Homo	(AF001628) interactor	protein AbiBr4 [Homo	sapiensj					nuclear autoantigen fo	14 kDa [Homo sapiens]	(AF072441) calcineurin binding	protein cabin 1 [Homo sapiens]	(AK001676) unnamed	protein product [Homo	alphal A-voltage-	dependent calcium	sapiens	SŴI/SNF complex 60 KDa subunit [Homo
		blastx.2	blastx.2	blastx.2							blastx.2		blastx.2		blastx.2		blastx.2			blastx.2
		350	351	352							353		354		355		356			357
		1005359	1005384	1005511		•					1005843		1005974		1006018		1006055			1006142
		HOELP29	HWLFG04	HYAAC49							HC00A71		HOUFB45		HIISTI14		HMCDB21			HSDEY08

403	377	773	418	172	699	1187	505	328	420	979		533	611
158	105	999	323	44	4	237	146	7		3	628	.0	201
47%	37%	27%	34%	%16	100%	27% 22%	82%	100%	%66	64%	41%	44%	40%
				gb AAD31938.1 AC0 07055 3	dbj BAA91430.1	gb AAC71813.1	gb AAA20645.1	gb AAD44480.1	emb CAB70899.1	gb AAC69438.1	-		
sapiens			-	(AC007055) unknown Homo sapiens]	(AK000927) unnamed protein product [Homo sapiens]	(AE001373) predicted secreted protein [Plasmodium falciparum]	light chain 3 subunit of microtubule-associated proteins 1A and 1B [Rattus norvegicus]	(AF078848) BUP [Homo sapiens]	(AL137735) hypothetical protein [Homo sapiens]	1(3)mbt protein	homolog [Homo	sapiens]	
				blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2			
				358	359	360	361	362	363	364			
				1006215	1006250	1006317	1006399	1006445	1006470	1006483			
				HOOJT92	HNORE03	HPCTS21	HFATL31	HE2JE77	HCBNX87	HNTDJ68			

840	221	840	614	294 428	515	136	215	17	235	395	438	488	571		908		449	1220	1294		
631	9	628	459	109 393	06	-	126	247	435	514	346	426	305	 }	8	}	75	473	1220		
%09	25%	49%	78%	33% 50%	%86	, le	73%	83%	70%	87%	740%	52%	10007	10078	27%	3	%66	7050	44% 44%		
	-			emb CAA30276.1	- H A A 36547 11	golyman	gb AAF24054.1 AF0 90942_1	oblA A B66528.11			17 0000 1	dbj BAA92090.1 	100000	gb[AAD40006.1 Ar1 50100_1	11 305304 4 4 11	gb AAA83/03.1	oblAAD34131.1lAF1	51894 1	emb CAB02879.1		
				heat shock protein e [Drosophila	十		942) PRO0657	SIL		polyprotein [inutiple sclerosis associated	retrovirus	(AK002129) unnamed protein product [Homo	sapiens]	(AF150100) small zinc finger-like protein	[Homo sapiens]	SpZ12-1 [Strongylocentrotus	purpuratus	protein [Homo sapiens]	predicted using Genefinder: Similarity	to Yeast mitochondrial	1 1 yk432a4.3 comes
				blastx.2		blastx.2	blastx.2	1	blastx.2	,		blastx.2		blastx.2		blastx.2		blastx.2	blastx.2		
				365		366	367		368			369		370		371		372	373		
				1006512		1006635	1006858	,	1006943			1006953		1007230		1007941		1007976	1008013		
				HTRAA05		HMTAL96	HDTJP21		HISEO81	,		НОДЕН02		HOCPP18		HODFP91		H6EDU06	HPCRD26		

374 blastx.2 (AF007791) secreted
XAG-2 homolog
375 blastx.2 PDI (E.C.5.3.4.1) [Bos taurus]
376 blastx.2 IEF SSP 9502 [Homo sapiens]
378 blastx.2 guanylate kinase [Homo sapiens]
379 blastx.2 bicaudal-D [Homo
sapiens
380 blastx.2 (AB005878) BYJ15
t
382 DIASIK.2 IDIN4-GO
383 blastx.2 pol gene protein; Xxx [Homo sapiens]
384 blastx.2 (AF019386) heparan
sulfate 3-0-
norms

	408	737	858		217	71	214	341		405	470	104	216	1106	1051	110	262	
	782 661	24	1		62	3	2	183	·	166	387	217	323	1282	1113	2	221	
	29%	93%	%/6		%99	%69	%59	%29		700%	65%	25%	55%	%5% 25%	54%	7000	42%	
	emb CAA48048.1	gb AAA35654.1	AbilB A A 13071 11	ابيد بمديعيطائمه	11.121.04 4.01.121		-1-1 A A OOO 20 11	gp AAAcoocoa		11 100000 11	gb AAA32067.1	114 4 113 4010 11 4 EO	gb AAF24010.1 ALO	dbj BAA91131.1		3/1//1200 2/20/2000	sp Q9Y6Y3 Cyang	
precursor [Homo	sapiens] T2 [Mus musculus]	liced	sapiens	matrix metalloproteinase, MT2MMP [Homo	╅	385) umamed product [Homo	sapiens	unknown protein [Homo sapiens]			Krueppel-related DNA- binding protein [Homo	sapiens	(AF090894) PRO0113	(AK000385) unnamed	protein product [Homo	sapiens	IDN4-GGTR14	PROTELLY.
	blastx.2	blastx.2		blastx.2		blastx.2		blastx.2			blastx.2		blastx.2	blastx.2			blastx.2	
	385	386		387		388.		390			391		392	393			394	
	1008326	1008327		1008332	_	1008335		1008337			1008345		1008349	1008350	2000		1008358	
	HBGBE52			нгнсв31		HPFDV51		HODEG13			HHPFK16		HARAL81	TTGT A T (1/2)	1 PC 1 PC - 1		HETDA81	

							_									
306	400 546	596 67	261	628	375	1273	1859	1404	1959	1279	85	692	296	1268		1151
	236 424	102	1	374	136	296	1404	.1270	1876	1145	7	296	m	174		375
45%	55% 51%	36%	\$2%	%56	51% 94%	100%	100%	%26	36%	30%	100%	79%	93%	100%		%26
gb AAC28019.1	dbj BAA91131.1	pir A26718 A26718	emb CAA16931.1	emb CAA11547.1		emb CAB54801.1					sp Q9Y6Y5 Q9Y6Y5	gb AAD53289.1 AF1	76818_1	gb AAD32538.1 AF1	05261_1	gb AAD27771.1 AF0
(AF062006) orphan G protein-coupled receptor HG38 [Homo sapiens]	(AK000385) unnamed protein product [Homo sapiens]	retrovirus-related pol polyprotein pseudogene - human 1	(AL021768) ATP binding protein-like [Arabidopsis thaliana]	(AJ223782) CDC10	[Mus musculus]	(AJ237734) ribophorin	II [Homo sapiens]				IDN4-GGTR14 PROTEIN.	(AF176818)	transcription factor AP-2 [Silurana tronicalis]	(AF105261) natural	killer cell receptor 2B4 [Homo sapiens]	(AF077038) unc-50
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2		blastx.2					blastx.2	blastx.2		blastx.2		blastx.2
395	397	398	399	400		401					402	403		404		405
1008359	1008379	1008396	1008401	1008406		1008423					1008429	1008445		1008764		1009017
HODE192	HODEG86	HODFA57	HODFB31	HTEMP79		HODEI59					НОДЕГ94	HPFCZ53		HWACN7	1	HOFMU69

					77038 1			
				related protein notinous			- 6	037
HETBR71	1009343	407	blastx.2	(AL049610)	emb CAB55700.1		187	° 66
			. •	(transcription				-
				elongation factor A				
	•			(SII)-like 1) [Homo				
-				sapiens]		7000	"	914
HE2ES17	1009349	408	blastx.2	CDV-1R protein [Mus	emb CAA71519.1	83%	ר-	
				musculus	1100000	040%	46	349
HELHM06	1009362	409	blastx.2	pre-mRNA splicing	gb AAA36649.1	9/60	3	· ·
				factor [Homo sapiens]	11 A COOK 24 11	100%	555	427
UEDESK1	1009388	411	blastx.2	(AC005034) gc-rich	gp AAC82330.1	2001		
TECESOT	2222			sequence dna-binding				
	-			factor [Homo sapiens]		7970	22	300
TENSOA?	1009398	412	blastx.2	protein Htf9C [Mus	emb CAA39515.1	/4%	5	2
1E23045	20001		****	musculus		7607	15.7	×
TT TT CITO	1000403	413	blastx.2	(AL031985)	emb CAB46381.1	%6/	171	•
HLHC10/	COT COOT	}		dJ228H13.1 (similar to				
				Ribosomal protein				
				[L21e] 1	11 V COOO 4 4 CH.	100%	55	231
UTTREAK	1009414	414	blastx.2	E1A-F [Homo sapiens]	4 dbj BAA02234.1	%09	449	622
HPFCV71	1009453	415	blastx.2	(AK002129) unnamed	dbjjbAA92090.11	3	:	
! !				protein product [round				
			1	Sapiens	dhilBA A05679.1	%88	127	423
HCABR46	1009479	416	blastx.2	BS1-2 Homo saprens	AbilB A A 23363.11	100%	96	479
HCRCB80	1009482	417	blastx.2	OTKZ/ [Homo	l		1	30,
200	1000033	418	hlastx.2	E2A/PRL fusion	gb AAA36764.1	99%	7	900
HPMLW/8	_	27.5						

				protein [Homo sapiens]				
HODJX27	1009854	420	blastx.2	(AJ005324) glutamate permease [synthetic construct]	emb CAA06471.1	%06	1	123
	<u> </u>							
HODJV85	1009857	421	blastx.2	ribosomal protein [Homo sapiens]	gb AAA36589.1	77% 35%	366 505	205 416
HOOJO38	1009918	422	blastx.2	ribosomal protein L14 [Homo sapiens]	dbj BAA13443.1	%86	П	603
HAOSI69	1009949	423	blastx.2	bcl-1 [Homo sapiens]	gb AAA58392.1	100%	271	1155
HSDIY67	1010157	424	blastx.2	hypoxanthine (guanine) phosphoribosyltransfer ase [Cricetulus longicaudatus]	emb CAA42198.1	%69	09	689
HSDZM65	1010299	425	blastx.2	(AJ010953) putative Ca2+-transporting ATPase [Homo sapiens]	emb CAA09425.1	100%	2	337
НРДОС39	1010400	426	blastx.2	(AB009282) cytochrome b5 [Homo sapiens]	dbj BAA23735.1	%66	171	608
HE2PS15	1010525	427	blastx.2	(AF054175) mitochondrial proteolipid 68MP homolog [Homo	gb AAC39909.1	100%	63	236
HTTKH22	1010596	428	blastx.2	weak similarity to	emb CAA86663.1	38%	887	1435

					_																			\neg
1561 949	1207	1274	1278	730	1630	1100	243	775	1603	1103	769	1612	1612	775	1600	1091	709	1085	1094	1097	727	1088	1085	1007
1442 692	290	1203	1246	236	1100	735		200	1100	753	203	1148	1160	284	1091	747	302	750	738	747	239	099	747	750
32% 27%	%86	%89	100%	82%	77%	82%	70%	23%	23%	34%	25%	73%	79%	27%	27%	31%	19%	27%	27%	30%	76%	27%	22%	33%
	gb[AAC39524.1]			dbilBAA19002.1																				
microtubule associated proteins; cDNA EST 1 1 EMBL: D37339 comes from this gene; cDNA EST FMRI : D6742	SAS [Homo sapiens]	[]		A7-1 Mins musculus												•						•		
	hlacty 2	- Creation		blacty 2	7:07:0																			·
	420	74		130) f																		<u>-</u>	
	1010608	1010020		1011000	1011030																			
	115101033	CCOCONTA		111111111111111111111111111111111111111	HFD V N40																			

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1331	1061	691	1106	730	730	249	938	246	673	206	273	1628	152	234	326	9		541	141	623		533	205		551	123	883
280	735	305	750	413	413	151	747	106	401	533	26	30	120	7	237	-		152	97	09		240	59		72	78	62
70%	23%	25%	22%	27%	78%	42%	32%	31%	23%	25%	24%	%66	54%	82%	73%	35%	_	%99	%99	%68		43%	44%		46%	43%	46%
												gb AAC50152.1		dbj BAA01393.1				emb CAA71771.1		emb CAA68868.1		emblCAA61914.1			emb CAA43408.1	-	gb AAC77439.1
												hypoxia-inducible	factor 1 alpha [Homo	2-oxoglutarate	dehydrogenase	precursor [Homo	sapiens	unnamed protein	product [Bos taurus]	coxsackie and	adenovirus receptor	interferon alpha /beta	receptor [Homo	sapiens]	GM2-activator protein	[Homo sapiens]	(AF039584) decay
												blastx.2		blastx.2		-		blastx.2		blastx.2		blastx.2			blastx.2		blastx.2
						·						431		432				433		434		435			436		437
						1						1011186		1011209				1011303		1011315		1011316		•	1011321		1011332
							•					HNORJ10		HPDRG92				HOFMQ81	,	HOFNF27		HOFND52			HOFNL96		HOFNF53

1706	784	1101	143	1249	582	736	004	605	552	020	2	1482		266	942	942	534	537	537	549	537
1155	191	880	66	377	88	602	2 5	576	124	500	+77	220	ì	114	202	235	229	229	202	232	235
41%	78%	32%	75%	%09	%98	%88	82%	%06 70%	100%	2000	93%	%000	2/66	100%	45%	45%	25%	46%	20%	46%	47%
	-			emb CAA08796.1	oblA A D 17527.11				dbj BAA01457.1		dbj BAA11829.1 	11 00011 4 4 61 61	dbj BAA11829.1	gb AAD27778.1 AF0 77045_1	emblCAA37624.1		•				
Total Catal	accelerating factor	One of the control of	CD35 [Kallus	embigin tus	norvegicus)	om	sapiens]		midkine [Homo	sapiens]	collagen binding protein 2 [Homo	sapiens	collagen binding protein 2 [Homo	(AF077045) ATP synthase epsilon chain	Homo sapicins	type A comagon [2003	raurus				
				blastx.2		blastx.2			Flacty 2		blastx.2		blastx.2	blastx.2		Diastx.2					
				438		439			940	}	441		442	443		444					
				1011360		1011499			1011627	/cc1101	1011607		1011608	1011755		1011773					
				HOFMU61		HOFOF35				HOLFJ3/	HOFME75		HOPKO74	HCBBA51		HDHEB13					

534	549	537	537	534	537	537	534	537	534	534	534	750	537	750	534	549	537	277	130	277	248	817		180	·
229	229	235	232	199	211	235	235	226	235	226	202	235	235	235	232	202	283	. 53	20	47	75	20		4	
45%	45%	49%	49%	45%	48%	46%	46%	47%	45%	44%	42%	35%	45%	34%	45%	40%	48%	32%	20%	31%	%16	100%		%6 <i>L</i>	
																			·	•	emb CAB71156.1	emb CAA91556.1		gb AAB34266.1	
																					(AJ271158) DAPIT protein [Rattus norvegicus]	CLC-7 chloride	sapiens]	scleraxis=basic helix-	loop-helix transcription factor [mice, embryos,
														,			,	-			blastx.2	blastx.2		blastx.2	٠,
																				-	445	446		447	
																					1011821	1011830		1011840	
									-												HDTMG36	HVVCJ38		HPAMY22	

			Peptide, 207 aa] [Mus				
	-		+	11. 1200114 11 A E1	100%	137	727
1011847	7 448	blastx.2	(AF161499) HSPC150 [Homo sapiens]	gb AAF29114.1 AF1 61499_1	0,001		COS
1011883	3 449	blastx.2	ke ca	gb AAF16704.1 AF1 17582_1	25%	7	760
1011901	1 450	blastx.2	sextal (AK000376) unnamed protein product [Homo	dbj BAA91124.1	94%	9	425
1011919	9 451	blastx.2	sapiens] (AL117452) hypothetical protein	emb CAB55934.1	%66	96	905
	1	Llogh 2	[Homo sapiens]	gb AAA03587.1	%16	125	1105
HPAMS93 1011948 HCRBN71 1011966	56 453		unknown [Saccharomyces	gb AAA79999.1	40%	94	024
HCRNC60 1012005	05 454	4 blastx.2	tropomyosin [Homo sapiens]	gb AAA36771.1	81% 91% 31%	769 1202 796	1236 1519 1245
					28% 27% 24%	766 766 925 1238	1257 1218 1212 1459
1012064	164 455	5 blastx.2	IFN-alpha responsive transcription factor	gb AAA58687.1	100%	392	925
HOVDZ22 1012136	36 457	57 blastx.2	unnamed protein	emb CAB69195.1	%68	2	82

1306 646	691	382 852	834	319	066	310	731	1716	300 443	
2 2	101	23 601	307	71	-	2	E .	. 64	452 550	
29%	%66	33% 27%	94%	%86	%16	%86	%86	%66	64% 41%	·
gb AAA80488.1	gb[AAA66351.1]	dbj BAA91635.1	emb[CAA25141.1]		gb AAA79184.1	emb CAA94861.1	emb CAA50574.1	gb AAA36458.1	gb AAF22026.1 AF1 18094_21	
100 kDa coactivator [Homo sapiens]	TB2 [Homo sapiens]	(AK001339) unnamed protein product [Homo sapiens]	DÇ classII	histocompatibility antigen alpha-chain [Homo sapiens]	DNA-PK [Homo sapiens]	inositol 1,4,5- trisphosphate receptor type 2 [Mus musculus]	peroxisomal acyl-CoA oxidase [Homo sapiens]	p78 protein [Homo sapiens]	(AF118082) PRO1902 [Homo sapiens]	
blastx.2	blastx.2	blastx.2	blastx.2		blastx.2	.blastx.2	blastx.2	blastx.2	blastx.2	
460	461	462	463		464	465	466	467	469	
1012454	1012459	1012469	1012482		1012519	1012536	1012542	1012545	1012599	
HCONM62 1012454	HOGEU69	нуувр93	HCOMV86		HCONJ23	HVVBL04	HVVAF65	HCONN76	норедээ	

1567	643	709	527	1099	.	250	2036 2856 1877 2252 2015 2078 2030 2021 1829 2018 1877 2039
89	101	2 648	48	44 1102		74	24 2032 141 123 222 135 117 117 117 117 69
100%	%86	93%	100%	%66 %66		100%	95% 97% 27% 23% 23% 22% 23% 22% 22% 22%
emb CAA30314.1	emb CAA53384.1	emb CAA55046.1	emb CAA62013.1	gb[AAD08640.1]		dbj BAA22054.1	emb CAA53052.1
 C1 inhibitor [Homo	protein kinase C mu	amiloride binding	NBK [Homo sapiens]	(AF026851)	assembly factor [Homo	(AB006202) cytochrome b small subunit of complex II	giantin [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	· .	blastx.2	blastx.2
470	472	473.	474	475		477	478
1012600	1012645	1012646	1012652	1012654		1012665	1012668
HCOOX52	HVVBK78	HVVBK73	HCOMW3	5 HVVBJ28		HVVAF41	HVVAS32

			_					_	_																		
1994	2039	2622	117	2700	2631	2628	2625	2616	2652	2538	2625	2634	2622	2694	2559	2715	2313	2610	2628	2565	2238	136	3910	3901	2616	3057	982
441	1281	2014	43	2437	2071	2125	2143	2128	1987	2032	2140	2206	2035	2053	. 1987	2104	2032	2065	2152	2032	2080	110	3779	3833	2035	2905	134
22%	20%	23%	84%	34%	22%	20%	24%	23%	23%	21%	24%	70%	23%	21%	24%	70%	25%	19%	19%	20%	78%	100%	22%	43%	21%	27%	100%
		<u> </u>						· .						~~~			•		-								gb AAB93872.1
																	•						•				voltage dependent anion channel form 3 [Homo sapiens]
							.,						-														blastx.2
																											479
								•						•		٠											1012678
												•															HVVAQ70

573	1990 2961	1155 1306 1375 1376	553	454	653	1605 203 1389 1407 1395 1605 1605 1653 1615 1615
79	302	1 1142 1310 1332	113	2	3	205 69 1186 1186 1195 1195 1375 1375 1490
%86 %86	%66 %66	99% 92% 54% 60%	48%	100%	%26	99% 100% 38% 34% 24% 26% 30%
gb AAB50715.1	gb AAA97878.1	gb AAA58391.1	emb CAA94699.1	emb CAB56526.1	dbj BAA91330.1	gb AAD56725.1
smooth muscle myosin light chain kinase, smMLCK [C-terminal]	specific 116-kDa vacuolar proton pump subunit [Homo sapiens]	receptor kinase [Homo sapiens]	agpet8 protein. [Schizosaccharomyces pombe]	(AL035593) dJ310J6.1 (novel protein) [Homo sapiens]	(AK000703) unnamed protein product [Homo sapiens]	(AF124727) acinusS [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
480	481	482	483	484	485	487
1012684	1012693	1012700	1012711	1013085	1013213	1013288
HVVAQ22	HTAES83	HOCMN67	HAPOW35	HOVJM48	HBJHY84	HFIAS44

Γ	T								_									_			_	_	
1207	538	675	3175	853	853	763	305	341	637	707	750	516	136	440	713		183	29	149	781	646	629	776
1112	269	43	2	. 7	53	70	n	6	281	633	649	139	2	165	54		317	159	181	2	7	7	·
40%	100%	%86	%66	34%	32%	27%	27%	25%	%86	%88	47%	%66	%06	40%	100%		%09	40%	63%	%09	%59	%65	23%
	gb AAB53427.1	gb AAD08657.1	gb AAC25585.1						dbj BAA91882.1			emb CAA35829.1		emb ÇAA30276.1	gb[AAA59581.1]		dbj BAA91205.1			gb AAA93261.1			
	non-histone chromosomal protein	(AF071172) HERC2	(AF053944) aortic	carboxypeptidase-like	protein ACLP [Homo	sapiens]			(AK001751) unnamed	protein product [Homo	sapiens	elongation factor 2	[Homo sapiens]	heat shock protein Drosophila melanogasterl	metalloproteinase	inhibitor precursor [Homo sapiens]	(AK000496) unnamed	protein product [Homo	sapiens]	zinc finger protein	C2H2-25 [Homo	sapiens]	•
	blastx.2	blastx.2	blastx.2			_			blastx.2			blastx.2		blastx.2	blastx.2		blastx.2			blastx.2	-		
	488	489	490						491			492		493	494		495			496			
	1013349	1013436	1013524						1013687			1013740		1013853	1014003		1014041			1,014204		•	
	HHAUD68	HEEAA89	HNTAK22						HOVBX78			HVVAW7	7	HSPSF84	HPAMV95		HOVBX22	_		HPDRB63		-	

99/	300	313	232	2467		1741	2434	2413	1624	890	2299		380	1061	352	167	480	794		158	112	48	1086	1700	109	1106	
47	-	200	1 7	L			1535		536	168	83		3	429	176	6	352	75		328	246	110	106	619	35	1083	
46%		46%	%99	\downarrow		%66	85%	36%	33%	100%	100%		100%	73%	77%	73%	44%	100%	2001	43%	54%	85%	100%	31%	 %96	100%	i
					emb CAA71575.1	11 4 4 4 5 8 6 8 3 1 1	Iricanocuravia			dbj BAA32209.1	11 00000 11	dbj BAA00338.1 	11 80598 V UTT	emolchargon 11	gb AAr138/2.1			10 4 10 4 DO	gb AAD49/45.1 AF1 72066_1	:: :: A A O1121 11	dbj BAA91131.1		dbilBAA90992.1				
		-				1		induced protein [Homo	sapiens]	(AB014888) MRJ	-	nthetase	Homo sapiens	SOX9 [Homo sapiens]	(AF180801)	peroxisomal long chain	acyl-CoA thioesterase	Ib [Mus musculus]	(AF172066) retinoic acid repressible protein	Homo sapiens	(AK000385) unnamed	protein product [Homo	sapiens	(ARU001/8) umanica	protein product [homo	sapiens]	
				,	blastx.2		blastx.2	-		blactx 2		blastx.2		blastx.2	blastx.2			•	blastx.2	•	blastx.2		١	blastx.2			
					497		498			80	}	200		501	502	ļ			504		505			206			
					1014252		1014432			1014405	1014403	1014646		1014730	1014754				1014869		1015010))		1015024			
					HNHG166		96Nddun			一	HPC1H41	HPCTV73		110000023	HCOrOss HOTERS	HOEBKSO			HOEER36		TIESON775	TESO CAL		HV ASTO6	22227711		

				_		_					_,												
215		615		319		9	543		61		364			1164	1204		971	1575	2428	1263	1736	602	681
3		1		5		136	136		298		125			934	41		159	1258	2063	991	1533	30	613
. 64%		%66	•	%19		40%	41%	-	%86		21%			%86	%66		33%	78%	30%	24%	30%	%56	26%
dbj BAA91118.1		emb CAB70908.1	•	dbj BAA91205.1	-	gb AAD35130.1 AE0	01691_4		dbj BAA35276.1		dbi BAA91131.1			sp O15532 SELW_H UMAN	gb AAA51597.1		emb CAB11184.1					dbj BAA91623.1	
(AK000370) unnamed	protein product [Homo sapiens]	(AL137756)	hypothetical protein [Homo sapiens]	(AK000496) unnamed	protein product [Homo	(AE001691) conserved	hypothetical protein	[Thermotoga maritima]	Lipoprotein RlpA	precursor. [Escherichia	(AK000385) unnamed	protein product [Homo	sapiens	SELENOPROTEIN W.	alcohol dehydrogenase	class III [Homo sapiens]	coronin-like protein	[Schizosaccharomyces	pombe]		-	(AK001322) unnamed	protein product [Homo
blastx.2		blastx.2		blastx.2		blastx.2			blastx.2		blastx.2			blastx.2	blastx.2		blastx.2					blastx.2	
202		208		605		510			511		512			513	514		515					516	
1015059		1015093		1015133		1015143			1015204		1015304			1015324	1015343		1015360					1015430	
HDPXP07		HE2KH02		HSPSI60		HOCOC26			HHEDC05		HODES86			HOPJUS7	HVCAC71	,	HSXCOSS					HLWDB78	

				sapiens]	- 500,000	%80	315	1232
HPCOU04	1015563	517	blastx.2	(AF093414) estrogen	gb AAC//43/.1	73%	1232	1390
				response element		77%	1309	1362
				binding protein		30%	282	518
				Saguinus Octubrasi	ahl A C 12944.11	%66	122	1054
HDPAT52	1015574	518	blastx.2	(AF036191) 1FA		42%	290	1054
				inducible protein	•	45%	467	1018
•	•		-	[Homo sapiens]		38%	458	1030
•		_				37%	467	1030
-			·			37%	473	1024
-						36%	473	1024
						34%	467	1024
					11 110047 111111111111111111111111111111	410%	139	549
HAOSL81	1015620	519	blastx.2	P24 protein [Mus	dbj BAA18947.1 	32%	815	970
				muscuius)	11 A POA 10 1 1 A B1	%66	232	1152
HIMWIU46	1015814	520	blastx.2	(AF148457) heterogeneous nuclear	gp AA57_1 48457_1			
				ribonucleoprotein,				
				alternate transcript				
		100	Llosty 2	(AC006486)	gb AAD11988.1	%06	3	1175
HSXCQ19	1015994	170	Diasta.2	PC85727 1 [Homo	-)	37%	880	1203
		٠		D-057/24_1 [110:10]		25%	30	515
				sapiens]		29%	595	867
						30%	15	299
						28%	63	284
						25%	523	88 88
	•					47%	341	397
HNOAG06	1016272	522	blastx.2	(AF001947) U4/U6-	gb AAC09069.1	100%	340	546
				associated KINA				

	T	Т.		т	1	T	γ	,	т
	089	1313	990	726	618	1196	1008	2063	2123
	24	69 55 1058	196	247	82	225	712	255	303
	75%	70% 65%	100%	73%	%66	100%	95%	%66	100%
	.gb AAF29566.1 AF0 95446 1	gb AAD11628.1	dbj BAA76400.1	gb AAF31432.1	dbj BAA91901.1	gb AAC16923.1	emb CAA62635.1	emb CAB51858.1	emb CAA33261.1
splicing factor [Homo sapiens]	(AF095446) syndesmos [Gallus gallus]	(AF060570) rig-1 protein [Mus musculus]	(AB016533) nuclear protein containing a WW domain (Npw38)	(AF213393) ATP- binding cassette protein [Mus musculus]	(AK001775) unnamed protein product [Homo sapiens]	(AF027299) protein 4.1-G [Homo sapiens]	seryl-tRNA synthetase [Homo sapiens]	(AJ132637) ATP-dependent metalloprotease YME1L [Homo sapiens]	precursor polypeptide (AA -21 to 782) [Homo
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	523	524	525	526	527	528	529	530	531
	1016351	1016740	1016768	1016939	1017051	1017227	1017374	1017461	1017565
	HE2KN09	нгурн58	HSPSB62	HE2FR37	нвлятизз	нсоод88	HNTSV21	HNORH33	HUFEF35

Homo sapiens 1 Homo sapiens 1 1017772 533 blastx.2 (AF092576) translation factor eIF3 p40 subumi; 1 p40 subumi	HTPCG30	1017694	532	blastx.2	(AF118078) PRO1848	gb AAF22022.1 AF1	%59	17	139
1017772 533 blastx.2 (AF092576) translation 1017801 534 blastx.2 (AF134404) delta-6 fatty acid desaturase Homo sapiens 1017825 535 blastx.2 (AF071081) proline-rich mucin homolog [Mycobacterium tuberculosis] 1018080 537 blastx.2 (AF161477) HSPC128 Homo sapiens 1018226 538 blastx.2 Zyxin [Homo sapiens] 1018243 539 blastx.2 Zyxin [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa sapiens] 1018459 540 blastx.2 (AF083385) 30kDa sapiens] Homo sapiens Homo sapiens		-	•	:	[Homo sapiens]	18094 17	45%	147	740
initiation factor eIF3 p40 subunit; 1 p40 subunit; 1 p40 subunit; 1 p40 subunit; 1 [Homo sapiens] 1017825 535 blastx.2 growth-regulating protein [Homo sapiens] protein [Homo sapiens] [Mycobacterium tuberculosis] protein product [Homo sapiens] p40 subunit; 1 [Homo sapiens] protein product [Homo sapiens] p40 subunit; 1 [Homo sapiens] p40 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens] [Homo sapiens]	HISBM03	1017772	533	blastx.2	(AF092576) translation	gb AAC84044.1	100%	751	924
1017801 534 blastx.2 (AF134404) delta-6 fatty acid desaturase Homo sapiens 1017825 535 blastx.2 growth-regulating protein [Homo sapiens 1018032 536 blastx.2 (AF071081) proline-rich mucin homolog [Mycobacterium tuberculosis] 101826 537 blastx.2 (AF161477) HSPC128 Homo sapiens 1018243 539 blastx.2 zyxin [Homo sapiens] 1018459 540 blastx.2 (AF000372) unnamed sapiens					initiation factor eIF3				
1017801 534 blastx.2 (AF134404) delta-6 fatty acid desaturase Homo sapiens 1017825 535 blastx.2 growth-regulating protein [Homo sapiens] 1018032 536 blastx.2 (AF071081) proline-rich mucin homolog [Mycobacterium tuberculosis] 1018080 537 blastx.2 (AF161477) HSPC128 Homo sapiens 1018243 539 blastx.2 zyxin [Homo sapiens] 1018459 540 blastx.2 (AF000372) unnamed sapiens 1018459 540 blastx.2 (AF083385) 30kDa sapiens Homo sapiens 1018459 1018459 1018459 1018450 1018					p40 subunit; 1	1	/000	-	532
1017825 535 blastx.2 growth-regulating	HOFAA79	1017801	534	blastx.2	(AF134404) delta-6	gb AAD31282.1 AF1	0200	+ · · ·	100
Homo sapiens 1017825					fatty acid desaturase	34404_1	83%	coc C	C 8 8
1017825 535 blastx.2 growth-regulating					Homo sapiens			18	
1018032 536 blastx.2 (AF071081) proline-rich mucin homolog [Mycobacterium tuberculosis] 1018080 537 blastx.2 (AF161477) HSPC128 Homo sapiens] 1018226 538 blastx.2 zyxin [Homo sapiens] 1018243 539 blastx.2 zyxin [Homo sapiens] sapiens 1018459 540 blastx.2 (AF083385) 30kDa sapiens Homo sap	HPRA 196	1017825	535	blastx.2	growth-regulating	gb AAA18898.1	100%	690	685
1018032 536 blastx.2 (AF071081) proline-rich mucin homolog [Mycobacterium tuberculosis] 1018080 537 blastx.2 (AF161477) HSPC128 1018226 538 blastx.2 zyxin [Homo sapiens] 1018243 539 blastx.2 zyxin [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa sapiens] 1018459 540 blastx.2 (AF083385) 30kDa sapiens]	_				protein [Homo sapiens]		2001	-	203
ich mucin homolog [Mycobacterium tuberculosis] 1018080 537 blastx.2 (AF161477) HSPC128 [Homo sapiens] 7 1018243 539 blastx.2 zyxin [Homo sapiens] 8 protein product [Homo sapiens] 1018459 540 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa sapiens] [Homo sapiens]	HRYEX71	1018032	536	blastx.2	(AF071081) proline-	gb AAD41594.1 AF0	32%		760
Mycobacterium tuberculosis	יייי איייין) } }			rich mucin homolog	71081_1	30%	_	633
1018080 537 blastx.2 (AF161477) HSPC128 [Homo sapiens] 1018226 538 blastx.2 zyxin [Homo sapiens] 1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa sapiens] Homo sapiens]					[Mycobacterium				
1018080 537 0lastx.2 (AF101777) 1151 CLEAR 1018226 538 blastx.2 zyxin [Homo sapiens] 1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]		0000	200	1.12.44. 2	(A E161/77) HQDC128	ohla A F 29092 1 A F 1	%66	23	745
1018243 539 blastx.2 zyxin [Homo sapiens] 1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]	HIMIVDD8	1018080	23/	DIASUX.2	(Ar 1014//) 1151 C129	61477 1			
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]	Ī	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	000	110000	myin [Homo saniens]	emblCAA64447.1	100%	143	1327
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]	OCHUVVH	1018220	000	Diasta.2	the charge of the control of the con		100%	1330	1857
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]							32%	1303	1455
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]							21%	662	1336
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens]					-		32%	134	346
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]				_			45%	1822	1914
1018243 539 blastx.2 (AK000372) unnamed protein product [Homo sapiens] 1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]							24%	5	295
1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]		1010010	625	Flooty 2	(AK000372) unnamed	dbi BAA91120.1	54%	194	45
Sapiens Sapi	HNNB157	1018243	757	Diasta.2	protein product [Homo		78%	355	230
1018459 540 blastx.2 (AF083385) 30kDa splicing factor; SPF 30 [Homo sapiens]					protein product		30%	363	244
Splicing factor; SPF 30 [Homo sapiens]	UCONTI 1	1018459	540	blastx.2	(AF083385) 30kDa	gb AAC64086.1	100%	186	668
[Homo sapiens]	TICATOOLI				splicing factor; SPF 30				
Linear CAL Linear CAL CALCOLO					[Homo sapiens]		, 000	20.	767
1018501 54 DIASIX.2 (ANDOOUTO)	HCOA WK	1018501	541	blastx.2	(AK000010) unnamed	dbj BAA90881.1	100%	104	430

1293	1455	1059	792	711	810	1204	2600	1800	2486	246	1237	1243	1183	1156	2588	1147	1195	1195	1201	1249	2531	1746	1791	1767	1198	1788	2588	1204	1237
1198	1381	409	418	394	418	332	1803	1204	1725	19	332	335	341	332	1803	332	332	332	332	299	1734	1204	1207	1219	335	1258	1794	491	335
100%	%96	%86	%95	47%	48%	%66	100%	100%	79%	63%	23%	23%	72%	23%	23%	22%	22%	25%	23%	23%	22%	23%		21%	22%	30%	22%	23%	19%
		emb CAA50897.1		•		emb[CAB05105.1]				•			-		-														
protein product [Homo	sapiens]	almost identical to	nRNP M protein,	acc.L03532 [Homo	sapiens]	dJ68O2.2 (myosin,	heavy polypeptide 9,	non-muscle) [Homo	sapiens	1															- ".				
	•	blastx.2		: .·	•	blastx.2														,									
		542				543																							
		1018772				1018802																							
∞		HVVCY25				HCOMB65														,									

1824	4016	470	1108	010	717	1800	1246	1773	1210	1791	1788	2435	1249	2477	1791	2471	2489	1785	1237	1794	2486	2480	1800	1872	1788	1776	1785	1800	2462	1800	
1291		1285	0 (727	230	1192	338	1219	353	1204	1219	1809	320	1806	1204	1806	1809	1207	332	1207	1881	1833	1207	1219	1195	1219	1339	1222	1806	1210	
		25%																										_			
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1788	1803	1767	1183	1473	1440	1800	228	1803	237	216	225	401	234	344	1305	387	294	295	131	349	128	380	861					945
1204	1237	1189	611	1228	1222	1714	29	1684	115	61	29	258	34	249	1192	121	118	149	n	284	42	282	151		<u>-</u> -			733
22%	23%	18%	25%	25%	29%	31%	35%	27%	31%	28%	792	24%	19%	28%	76%	23%	61%	%69	62%	%65	23%	47%	%66					42%
				-			•		-				•			gb AAC08737.1							gb AAC12956.1					emb CAA19575.1
					1											(AF010144) neuronal	thread protein AD7c-	NTP [Homo sapiens]					(AC004537) similar to	tumor suppressor	AF044076	(PID:g2829208)	[Homo sapiens]	(AL023859) tma-
				•												blastx.2							blastx.2					blastx.2
																544							545					546
																1018907		•					1018943			•		1019326
															·	HVVCF30							HODBV21		•			HCHIMD81

423		1	1192		607		1	2/2		•		919		468			830		(1	731						751	
85			110		314	71 648		CT				134	210	103			234		576	72		88	298	1146	1871	.131	
73%			100%		41%	54%	2/2/	%66				100%	2270	100%			%86 		48%	34%	20%	63%	%66	78%	42%	. 100%	
			emb CAA42641.1		dbj BAA25253.1		. 07077	emb CAB44749.1				gb AAF29106.1 AF1	01491 1	gb AAC05810.1			dbj BAA91259.1		emb CAA42610.1				emb[CAB70844.1]	-		gb AAA59930.1	
splicing endonuclease	subunit [Schizosaccharomyces	pombe	phosphate carrier	protein [Homo sapiens]	(AB012223) ORF2	[Canis familiaris]		(AL035494)	dJ635G19.2.1 (novel	protein (isoform 1))	[Homo sapiens]	(AF161491) HSPC142	Homo sapiens	(AC002394) Unknown	gene product [Homo	sapiens	(AK000566) unnamed	protein product [Homo	zine finger protein	Rattus norveoicus	[marga ron cannay]		(AT.137618)	himothetical protein	Itypometreal protein	CCAAT-box DNA	binding protein subunit
			blastx.2		blastx.2			blastx.2				blastx.2		blastx.2			blastx.2		hlacty 2	7.v.cpro			hlasty 2	-		blastx.2	
			547		548			549				550		551		_	554		444	3			556	3		557	
			1019338		1019409			1019585				1019608		1019749			1019892		1010017	1013342			100000	100001	•	1020130	227227
			HAZAR95		HMWFS51			HE8SD82				HSPSN08		HVCAH24			HJPCF71		355 A 5777	hvCAE/0			11000011	+111MOOU		HPDR703	

			· -	NF-YB [Homo sapiens]				
HOCPO73	1020180	828	blastx.2	protein p84 [Homo	gb AAA53571.1	93%	3646	2024
				sapiens]	-	%26	3881	3636
						100%	3993	3940
			_			45%	1832	1758
	`					37%	1888	1841
HNKDT10	1020832	529	blastx.2	(AF191018) E2IG3	gb AAF09482.1 AF1	%66	11	436
				[Homo sapiens]	91018 1	93%	426	569
HWHGO2 5	1020841	260	blastx.2	(AF090942) PRO0657 [Homo sapiens]	gb AAF24054.1 AF0 90942 1	%89	·C	137
HWMNE3	1020852	561	blastx.2	(AF083384) 45kDa	gb AAC64085.1	%86	169	675
				splicing factor; SPF 45				
HUSY175	1020878	562	blastx.2	(AJ242540)	emb CAB62280.1	33%	19	684
				hydroxyproline-rich		35%	29	624
				glycoprotein DZ-		34%	67	624
				HRGP [Volvox carteri		32%	129	684
				f. nagariensis]		34%	49	624
•	•					32%	29	714
				,		31%	19	969
						32%	29	624
	-		•		,	73%	64	714
				,		31%	29	654
						30%	64	654
						78%	64	969
					<u> </u>	34%	184	624
	-					31%	184	783
HSDFS07	1020904	263	blastx.2	(AL033534)	emb CAA22127.1	27%	186	118
				nypometical serme-men	<u>····</u>			

	305	1649		937	1227	614		230		0,70	007		689	3		3	220			1327				
	183	036	200	20	949	3	·	159		90,	479		210	717		209				890				
	%56	1,000%	10078	%68	89%	85%		100%			75%		2000	30%		100%		%09 —	36%	45% 28%	24%	30%	38%	
	emb CAB55759.1		emb[CAA80599.1]	-11 A PE34240 11AFO	goldar 34270.1 m. c 75704_1	A A IT SOOD TIA A	gb AAD18083.1 AA D18083	gb[AAC59341.1]			dbj BAA91205.1	.		emb CAB66159.1		114 4 4 70014 11	gb AAA/9014.1	dbj BAA91298.1						
[Schizosaccharomyces	+	[Chlorocebus aethiops]	hnRNP G protein	Homo sapiens	(AF075704) neuronal glutamine transporter	Rattus 1	(AF129756) G4 [Homo	(A E038616) email	tumor anitgen t-ag	[Simian virus 40]	(AK000496) unnamed	protein product [Homo	sapiens	(AJ011376)	hypothetical protein	Homo sapiens	ORF_0109 Fecherichia colil	(AK000633) unnamed	protein product [Homo	sapiens]				
	blastx.2	- 	blastx.2		blastx.2		blastx.2		blastx.2		Lloots 2	Diasta.4		blastx.2			blastx.2	hlastx 2	- Oleans				•	
	564		565		995		292		268		1			570			571	577	7/6					
	1021208		1021323		1021327		1021661		1021666			1021682		1021759	701701		1021794	1000010	1022018					
	HCOPC09		HVVBT22		HAMFW6		HPMBW8	9	HOFMK02			HOUHK71		C/X/OODIT	HC00 W 2	1	HACMT02		HISDS67					

						٠,																						
1372	780	1193	1319	2421	1585	1770	2211	1506	1917	1966	1536	1813	1063	2440	3	840	435	222		836	1570	445	95		1072		247	
1232	2000	1143	252	1765	1316	1573	1744	1417	1744	1736	1447	1718	. 38	2402		445	115	43		m	839	101	33		176		2	
%42	48%	41%	%65	33%	31%	39%	21%	46%	73%	27%	38%	36%	%16	95%		46%	43%	23%		100%	100%	%26	100%		%66		%001	
			oblA A C 00620.11		,							•	dbj BAA91548.1			gb AAA62280.1	-			gb AAD01548.1		gb[AAC31609.1]			emb CAB09725.1		emb CAB43242.1	
-			(AC002291) Similar	ATP-dependent RNA	Helicase [Arabidopsis	thaliana							(AK001197) unnamed	protein product [Homo	sapiens	epidermal growth	factor receptor kinase	substrate [Homo	sapiens	(AF015040) NUMB	protein [Homo sapiens]	(AF081280)	nucleoplasmin-3	[Homo sapiens]	ribonuclease HI large	subunit [Homo sapiens]	(AL050022)	hypothetical protein
			hlachy 7										blastx.2			blastx.2				blastx.2		blastx.2			blastx.2		blastx.2	
			573	3									574			575				978		577			578		280	
			100001	1007701									1022059			1022082				1022162		1022167			1022313		1022663	
			HOVA 459	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\									HOCPL72			HCOMIM0	S			HLYDC86		HTLEP21			HVCAS77		HSKJP93	

				[Homo comions]				
				Troing sapiens		/007	700	305
HAZAA64	1022719	581	blastx.2	predicted using	emb CAA99938.1	43%	304	<u></u>
				Genefinder; Similarity				
				to Prototheca 1 1 gene;				-
				cDNA EST yk386c1.3				
				comes from this gene;	•			
		_		cDNA EST yk38				
HSEBD72	1022904	582	blastx.2	(AF044773) breakpoint	gb[AAC08964.1]	%68	164	280
				cluster region protein 1				
				[Homo sapiens]				
HDPYE27	1022911	583	blastx.2	(AF062346) zinc finger	gb AAC42601.1	100%	428	1066
				protein 216 splice				
				variant 1 [Homo				
				sapiens				
HCROL51	1022997	584	blastx.2	PEG1/MEST [Homo	emb CAA72297.1	100%	20	1030
				sapiens]				
HDTCC55	1023046	585	blastx.2	(AL117483)	emb CAB55956.1	33%	306	533
				hypothetical protein		35%	4	105
				[Homo sapiens]		42%	105	182
HOOJ091	1023049	586	blastx.2	envelope protein	gb AAA88027.1	%69	m	230
-				[Homo sapiens]				
HCE4K28	1023227	587	blastx.2	(AF021792) Bcl-	gb AAB72092.1	- %86	262	750
				X/Bcl-2 binding protein				-
				[Homo sapiens]	-			
HPDVY62	1023264	588	blastx.2	DOCK180 protein	dbj BAA09454.1	%66	7	1126
				[Homo sapiens]				
HLHDK42	1023339	589	blastx.2	modifier 2 [Mus	emb CAA40012.1	61%	28	300
				musculus]		85%	3	62
HL.DRO55	1023375	590	blastx.2	coiled-coil like protein	gb AAB61902.1	%86	5	388
N. S.								

(AF064603) GA17 protein [Homo sapiens]
(AK001103) unnamed
protein product [Homo
sapiens
(AF104670) cell cycle
protein [Homo sapiens
chaperonin-like protein
Homo sapiens
ribosomal protein L34
Homo sapiens
ribosomal protein L34
TO T
INCO protein protein
(A FOOK264)
102000 LC)
recombination and
sister chromatid
cohesion protein
homolog [Homo
sapiens
1
vacuolar H+ ATPase E
subunit Homo sapiens

		blastx.2	(AL 109978) hypothetical protein				
7		1.10.ctv 7		dbj BAA20128.1	%16	3	392
1024472	700	7.VJSBIQ		11 1200101 111	%26	902	1231
1024556	603	blastx.2		gb AAB183/4.1 :	57%	198	275
1024624	604	blastx.2	(AF006084) p41-Arc	gb AAB64189.1	%99 %99	1029	1112
1024915	909	blastx.2	(AL050273) hypothetical protein	emb CAB43374.1	100%	203	710
			[Homo sapiens]	110 4 10 4 50 5 11	100%	48	971
1025047	909	blastx.2	(AJ249366) epsilon- COP protein [Homo	emolcAbbbbccc			
			sapiens	LIC 4 021052 11	39%	7	403
1025102	209	blastx.2	(AL031640) /prediction=(method:""	emologisch			
			genscan"", version:""1.0"", 1.1.1				
			target:SPTREMBL::06				
			087	-	%96	89	1225
1025231	809	blastx.2	(AB015597) hTIM1 Homo sapiens]	dbj BAA30499.1	797	130	c
1025327	609	blastx.2	(AK000385) unnamed protein product [Homo	dbj BAA91131.1 	04%	7	
			sapiens	11 A D 1 1850 11 A F 1	%86	183	512
1025359	.610	blastx.2	(AF110776) adrenal gland protein AD-003	10776_1	100%	512	766

1025366	611 blastx.2	F35C11.4	emb CAA90244.1	30%	162	632
		[Caenorhabditis elegans]		31%	743	1060
612 blastx.2	7	(AF083242) HSPC024- iso [Homo sapiens]	gb AAD39840.1	%06 %88	313	387
613 blastx.2	7	TRAM protein [Homo sapiens]	emb CAA45218.1	100%	175	1296
614 blastx.2	1~	retinoblastoma-binding protein mRbAp48 [Mus musculus]	gb AAC52275.1	96% 40%	124	1473
615 blastx.2	1 ~	GTP-binding protein (rab2) [Canis familiaris]	gb AAA30888.1	100%	444	719
616 blastx.2	١.,	(AF055010) unknown [Homo sapiens]	gb AAC09360.1	99%	1567 1337	2760 1594
617 blastx.2		follistatin-related protein FLRG [Homo sapiens]	gb AAC64321.1	100%	36	824
619 blastx.2	1	(AF054284) spliceosomal protein SAP 155 [Homo sapiens]	gb AAC97189.1	100%	123	497
620 blastx.2		(AK001093) unnamed protein product [Homo sapiens]	dbj BAA91500.1	100% 100% 31% 36%	1913 1654 1994 2405	2632 1848 2689 2755
621 blastx		putative glycosyl transferase [Schizosaccharomyces	emb CAB10854.1	55% 76% 42% 40%	488 389 272 92	616 439 376 202

524	378 566	1565	241 975 63	277	393	462	1004	667
54	557 601	9	35 865 7	275 74	22	91	123	2
74%	51% 66%	%66	78% 94% 52%	100%	100%	100%	73%	
gb AAD34062.1 AF1	gb AAF29584.1 AF1 13685 1	gb AAF29040.1 AF1 61553 1	emb CAA48726.1	gb AAC72409.1	emb CAB37641.1	emb CAB37641.1	gb AAD27851.1 AF1 32552_1	gb AAB37433.1
-	(AF113685) PRO0974	3PC068	ionine no	(AF102265) N-acetylglucosamine-phosphate mutase	(AL031663) d1461P17.6 (Major Epididymis-specific	(AL031663) dJ461P17.6 (Major Epididymis-specific	(AF13252) BcDNA.GM01838 [Drosophila	BB1=malignant cell expression-enhanced gene/tumor 1 line, Peptide, 342 aa] [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
622	623	624	625	626	627	628	630	631
1026913	1026979	1027007	1027207	1027269	1027484	1027486	1029191	1029484
HMUAQ0	5 HCOMA45	HE2K145	99ОДДОН	HVVAT45	HVCAG18	HPTXK72	HDTLR06	HPAMG11

1176	1516	1429	1899	1881	640	130	2158	96	462	2271	423	2696	1092
52	1097	1094	1747	1735	999	74	95	1	349	52	259	279	22
100%	32%	32%	45%	34%	48%	36%	%66 ·	%96	%98	%66	%59	100%	%66
emb CAB70786.1	emb CAA46283.1						dbj BAA07552.1		dbj BAA85182.1	emb CAA49992.1	gb AAA31142.1	gb AAC07984.1	dbj BAA90894.1
(AL137520) hypothetical protein [Homo sapiens]	extensin [Volvox carteri]	÷	-				ha1025 is new [Homo	sapiens]	(AB033168) nuclear protein ZAP [Mus musculus]	member of DEAD box protein family [Homo sapiens]	valosin-containing protein [Sus scrofa]	(AC004472) TERA_HUMAN [Homo sapiens]	(AK000031) unnamed protein product [Homo sapiens]
blastx.2	blastx.2						blastx.2		blastx.2	blastx.2	blastx:2	blastx.2	blastx.2
. 632	633						634		929	637	638	639	640
1030870	1030871						1030963	-	1031085	1031316	1031328	1031329	1031435
HSDJR27	HOCPL33				-		HPCQN80		HODFZ16	HWGAE28	HOFNZ21	ноокн25	HDTIL75

1751	631	<u> </u>		1328			1204			672			785				400		431		ļ	801	889
6	512			09	3		365			379			105				. 260		48			700	633
%66	100%			%00	27.66		41%			100%	2		%86				%26	•	1000%	100.78		91%	100%
gb AAF01278.1	gb AAD17301.1			111 110 CO 20 CT 1 1 1 EO	gb[AAD258/0.1 AF0	- 10107 - 10107	mblC 4 B06436 11	l::octoody)one		11001024 4011	emb[CAA00100.1]		-LIA A CSO419 11	golaracourts:1			dbi BAA91040.1		11 0001 5.	emb CAB41269.1		gb AAB72234.1	-
ivating tor 1	+		DNA glycosylase	1		chain family member	mulb inouto saprema	hypothetical protein Rv0712	Mycobacterium	tuberculosis	S100 calcium-binding	protein A13 (5100A13)	HOIIIO Sapicius	putative T1/S12	receptor binding	protein precursor	(AVOOOSA) unnamed	protein product [Homo	sapiens	(AL049705)	hypomeneal protein	novel ORF [Homo	sapiens]
blastx.2	blacty 2				blastx.2			blastx.2			blastx.2			blastx.2				DIASTX.2		blastx.2		11.1	Diastx.2
641	247				643			644			645			646				647		648		-	049
1031451	2021501	1031600			1031922			1031988			1032475			1033653				1034320		1034471			1034539
HPRSB55	$-\dagger$	HPDPY36			HETF147			HMAMI21			HC00085	, ,		HE2DO62				HEBAE89		HKDBF43			HVVCT43

[_	9	150	T	T	lia -		T	·
841	986	115	974	1299	435		786	357
806	693	468	93	439			286	73
91%	48%	38%	%66	100%	%66	:	30%	32%
	gb AAF19993.1 AF2 14634_1	gb AAA88036.1	emb CAB60141.1	gb AAB69324.1	dbj BAA91872.1		gb AAD30426.1 AF1 17723_1	emb CAB01127.1
	(AF214634) polyA binding protein [Homo sapiens]	unknown protein [Homo sapiens]	(AJ246001) spastin protein [Homo saniens]	(AF013249) leukocyte- associated Ig-like receptor-1 [Homo 1	(AK001738) unnamed protein product [Homo sapiens]		(AF117723) seed maturation protein PM27 [Glycine max]	predicted using Genefinder; Similarity to Drosophila RNA 1 this gene [Caenorhabditis
ļ	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2		blastx.2	blastx.2
(1)	000	651	652	653	655		929	657
1025425	1035435	1035602	1035988	1036583	1036973		1037108	1037131
UTTIDDO	שנאתווה	HODG046	HNTTB23	HVVBV73	HTEPV42	·	HFTCG52	HKAOB40

1037438 659 blastx.2 GAP-associated protein pirlA38218 A38218 74% 56% 681 8 63% 681 681 680 blastx.2 GAP-associated protein gb AAB49682.1 100% 150 41% 43 1 1038344 661 blastx.2 transferase Homo sapiens 1038378 662 blastx.2 transferase Homo sapiens 1038718 664 blastx.2 transferase Homo sapiens 1038718 665 blastx.2 transferase Homo sapiens 1038718 665 blastx.2 transferase Homo sapiens 1038718 665 blastx.2 transferase Homo sapiens 1038915 665 blastx.2 transferase Homo sapiens 1038917 666 blastx.2 transferase Homo sapiens 1038918 666 blastx.2 transferase Homo sapiens 1038918 666 blastx.2 transferase Homo sapiens 1039017 666 blastx.2 transferase Homo sapiens 100% 114 114 115 1039017 666 blastx.2 transferase Homo sapiens 100% 114 115 1039017 666 blastx.2 transferase Homo sapiens 100% 114 115 1039017 666 blastx.2 transferase Homo sapiens 100% 114 115 1039017 666 blastx.2 transferase Homo sapiens 100% 114 115 115	HPDVE37	1037303	658	blastx.2	n C [Homo	emb CAA60671.1	100%	118	880
1037438 659 blastx.2 GAP-associated protein pultatorated p					sapiens	15 A 287181A 38718	74%	50	811
1037480 660 blastx.2 ubiquitin-like protein gb AAB49682.1 100% 150 1037480 660 blastx.2 Bos taurus timase Edomo sapiens sapiens 1038713 662 blastx.2 transferase [Homo sapiens] t	T.FS74	1037438	629	blastx.2	GAP-associated protein	Disposion social ind	26%	681	875
1037480 660 blastx.2 ubiquitin-like protein gb AAB49682.1 100% 150					p190 - rat		63%	43	152
1037480 660 blastx.2 ubiquitin-like protein golAADA7002.11 1038344 661 blastx.2 kinase [Homo sapiens] nucleoside-diphosphate emb CAA68877.1 99% 44 1038378 662 blastx.2 candothelial cell growth gb AAA60043.1 100% 114 11038717 663 blastx.2 glutathione S- transferase [Homo sapiens] gb AAA35934.1 100% 110 1038718 664 blastx.2 glutathione S- transferase [Homo sapiens] gb AAA35934.1 100% 114 1038915 665 blastx.2 (AL080156) hypothetical protein Homo sapiens] Homo sapiens] Homo sapiens] Homo sapiens] 1039017 666 blastx.2 beta-spectrin [Homo gb AAA60580.1 20% 15 16 16 16 16 16 16 16						11 4 5 4 5 4 6 6 8 2 1 1	100%	150	434
1038344 661 blastx.2 mucleoside-diphosphate emb CAA68877.1 99% 44	ORG51	1037480	099	blastx.2	ubiquitin-like protein	gb AAB49082.1	2,001		
1038378 662 blastx.2 factor [Homo sapiens] 1038378 662 blastx.2 factor [Homo sapiens] 1038717 663 blastx.2 glutathione S- transferase [Homo sapiens] 1038718 664 blastx.2 glutathione S- transferase [Homo sapiens] 1038915 665 blastx.2 (AL080156) emb CAB45747.1 94% 415 1039017 666 blastx.2 beta-spectrin [Homo sapiens] 1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31 104 1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31 100% 31 100% 104 104 104 104 100% 104 100% 104 100% 104 100% 104 100% 104 100% 104 100% 104 100% 104 100% 104 100% 100% 104 100%	5	1000044	199	blastx.2	nucleoside-diphosphate	emb CAA68877.1	%66	44	520
1038378 662 blastx.2 endothelial cell growth gb AAA60043.1 100% 234 1038717 663 blastx.2 glutathione S- transferase [Homo sapiens] 1038718 664 blastx.2 glutathione S- transferase [Homo sapiens] 1038718 665 blastx.2 (AL080156) emb CAB45747.1 94% 415 1039017 666 blastx.2 beta-spectrin [Homo gb AAA60580.1 20% 1 1039290 668 blastx.2 TRAF4-associated 104 1 1039290 668 blastx.2 TRAF4-associated 104 1 1039290 1039290 1068 blastx.2 TRAF4-associated 104 1 10388015 100% 100% 100% 1 1039290 1039290 104 100% 100% 1 10388015 100% 100	AMM/	1020244	3		kinase [Homo sapiens]		10007	114	1559
1038717 663 blastx.2 glutathione S- gb AAA35934.1 100% 234 1038717 663 blastx.2 glutathione S- gb AAA35934.1 100% 114 1038718 664 blastx.2 glutathione S- gb AAA35934.1 100% 114 1038915 665 blastx.2 (AL080156) hypothetical protein Homo sapiens Homo sapiens Gb AAA60580.1 99% 1 1039017 666 blastx.2 beta-spectrin [Homo gb AAA60580.1 20% 16 1039290 668 blastx.2 TRAFF4-associated gb AAD24202.1 U83 100% 31 1039290 668 blastx.2 TRAFF4-associated gb AAD24202.1 U83 100% 31 103820 103820 104 104 104 104 1038218 100% 104 104 104 1038218 100% 100% 104 104 1038218 100% 100% 100% 104 1038218 100% 10	77AT03	1038378	662	blastx.2	endothelial cell growth	gb[AAA60043.1]	100%		, , , , , , , , , , , , , , , , , , ,
1038717 663 blastx.2 glutathione S- gb/AAA35934.1 100% 110					factor Homo sapiens	AND D 235034 11	100%	234	356
1038718 664 blastx.2 glutathione S- transferase [Homo sapiens] 1038915 665 blastx.2 (AL.080156) 1038915 665 blastx.2 (AL.080156) 1039017 666 blastx.2 capiens 1039017 666 blastx.2 capiens 1039290 668 blastx.2 capiens 1039290 668 blastx.2 capiens 1039290 668 blastx.2 capiens capien	OOMA8	1038717	663	blastx.2	glutathione S- transferase [Homo	i contrologi	100%	110	235
1038718 664 blastx.2 glutathione S- gol/AAA.2023.1.1 Sapiens Emb CAB45747.1 94% 415 1038915 665 blastx.2 (AL080156) hypothetical protein [Homo sapiens] Ehomo sapiens Homo sapiens Ehomo sapiens Ehom			-		sapiens	1114 4 4 2 5 0 3 4 11	100%	114	578
1038915 665 blastx.2 (AL080156) emb CAB45747.1 94% 415	VCCQ82		664	blastx.2	glutathione S- transferase [Homo	go AAA33371.1)))		
1038915 665 blastx.2 (AL080156) emb CAB43/47.1 770					sapiens	11 00000	7010	415	525
Homo sapiens Homo sapiens 1039017 666 blastx.2 beta-spectrin [Homo gb AAA60580.1 99% 1 20% 7 21% 28 22% 16 19% 1 19% 1 19% 1 19% 1 19% 1 1039290 668 blastx.2 TRAF4-associated 194 1 194 1 1039290 668 blastx.2 TRAF4-associated 194 1 194 1 1039290 100% 31 100% 31 100% 104 1 104	TGFP54	1038915	999	blastx.2	(AL080156) hypothetical protein	emb CAB45/4/.1	R.		
1039017 666 blastx.2 beta-spectrin [Homo gol.Arx.00.2001.] 20% 7 21% 28 22% 16 19% 1 19% 1 19% 1 19% 1 1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31 1039290 668 blastx.2 TRAF4-associated 194 1 104 1					Homo sapiens	11 A A 60580 11	%66	-1	2172
21% 28 22% 16 19% 1 22% 16 19% 21 1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31	RODF41	†-	999	blastx.2	beta-spectrin [Homo	goldenormaniag	70%	7	2091
1039290 668 blasts.2 TRAF4-associated gb AAD24202.1 U83 100% 31	! 				sapiens		21%	28	2151
1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31							22%	16	2187
1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31							19%	 1	2118
1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31							22%	184	2193
1039290 668 blastx.2 TRAF4-associated gb AAD24202.1 U83 100% 31							%09	2127	2351
	PAMC6	0 1039290	899	blastx.2	TRAF4-associated	gb AAD24202.1 U83 194 1	100%	31	1242

				sapiens]				
HLMHM8 3	1039491	699	blastx.2	(AK001123) unnamed protein product [Homo	dbj BAA91513.1	44%	158	616
				sapiens]				
HOUDK70	1039538	029	blastx.2	(AF091083) unknown [Homo sapiens]	gb AAC72952.1	100%	285	1142
HMEFK29	1039652	672	blastx.2	(AC002398) F25965_1 [Homo sapiens]	gb AAB81199.1	54% 100%	808	993
HAOSK79	1039663	673	blastx.2	(AJ223953) hPTTG [Homo sapiens]	emb CAA11683.1	100%	54	629
HPDRV42	1039689	674	blastx.2	P58 [Homo sapiens]	gb[AAC50331.1]	100%	135	1649
HOPJD35	1039703	675	blastx.2	(AF035262) BAF57 [Homo sapiens]	gb AAC04509.1	100%	121	1353
HTFNP84	1039748	9.29	blastx.2	ect2 [Mus musculus]	gb AAA37536.1	94%	73 27	1227 125
нтнрт76	1039871	229	blastx.2	(AL117404) hypothetical protein Homo sapiens	emb CAB55905.1	%86	7	564
HSYEC21.	1039891 ·	678	blastx.2	adenosine triphosphatase [Homo	gb AAA35999.1	99% 37%	432 1 3819	2372 435 3890
HKGC025	1040384	619	blastx.2	NAP [Homo sapiens]	gb AAC37544.1	%66	3	665
HNOJN70	1040385	089	blastx.2	(AF062594) nucleosome assembly protein [Rattus norvegicus]	gb AAC67388.1	100%	297	425
HADFS31	1040388	681	blastx.2	(AF214680) C3HC4- like zinc finger protein [Homo sapiens]	gb AAF30180.1	88% 92%	449 145	832 504

1040569	682	blastx.2	nonhepatic arginase	dbj BAA13138.1	97%	651	1175
	69	Flooty 2	Homo sapiens	dbj BAA91413.1	47%	303	557
1040591	680	Diasta:	protein product [Homo	:			
			sapiens		63%	84	527
1040620	684	þlastx.2	thioesterase II [Homo sapiens]	eno CA-A0002-1.1	7910	777	687
1040631	685	blastx.2	zinc-finger helicase	gb[AAC39923.1]	9170	117	3 6
1040694	989	blastx.2	tumor susceptibility	gb AAC52083.1	100%	0110	175
1040826	687	blastx.2	(AK001550) unnamed protein product [Homo	dbj BAA91751.1	63% 75% 31%	108	938 398 618
			sapiens	AND A 642232 11	%96	355	999
1040913	889	blastx.2	protein of unknown function [Homo			,	
			sapiens	1-1 A D 81 40 £ 11	%90	20	415
1040925	689	blastx.2	(AF005855) anon2A5 [Drosophila]	gb AAB61480.1			
			melanogaster	1100 100 11	100%	539	1384
1040932	069	blastx.2	protein antigen [synthetic construct]	emb CAA01162.1	%55	226	474
1041049	691	blastx.2	bcn92 [Drosophila subobscura]	emolcAb55511:1	7970	7	356
1041070	692	blastx.2	(AF161479) HSPC130	gb AAF29094.1 AF1 61479 1	74%	c	
1041900	693	blastx.2	NF-AT3 gene product	gb AAA79175.1	100%	33	212
			FROMO Saprova		39%	421	612

586	3432	338	2256	637	 494	852	870	48	51	138	077	600	629	900 900	629
455	259	174	2717	188	 180	52	301	632	584	632	070	74/	790	2 6	/90
40%	%001	%59	%56	40%	 100%	100%	100%	31%	34%	33%	52%	0,70	36%	3,600	39%
	gb AAA61246.1	gb AAD15797.1	dbj BAA03074.1	gb AAA76738.1	emb CAA54687.1	emb CAA30678.1	dbj BAA19122.1	pir A24264 A24264							
	ubiquitin-activating enzyme E1 [Homo sapiens]	(AF055470) ZNF258 [Homo sapiens]	Mel-18 protein [Homo sapiens]	unknown [Homo sapiens]	ATL-derived factor/thioredoxin [Homo sapiens]	pre-pump-1 proteinase (AA -17 to 250) [Homo sapiens]	(AB000468) zinc finger protein [Homo sapiens]	proline-rich protein	MP2 - mouse	(fragment)		•			
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	-						
	694	969	969	869	669	701	702	703	•						
	1042462	1042649	1042859	1042951	1043273	1043532	1043553	1044199				<u>-</u>			
	HSPSI42	НРDQD23	HWMBA1 0	HETKL27	HPCTH04	HVVBM94	HVVBC43	HTPDM31				-			

662	662	439	93		834			968			1738	493		1657	1319		1269	932	260	614	593		855		1067	1174	558		
790	790	53	-		424			ന			749	20		1316	1101		940	465	9	315	450		160		309	1067	34		
41%	38%	%68	48%		72%			77%			45%	47%		%89	73%		77%	41%	25%	71%	31%		100%		100%	%26	%96		
		dbj BAA05923.1			gb[AAD17989.1]			gb AAB17729.1		,	gb AAC14192.1			gb AAF31432.1			gb AAB00699.1	•					gb AAC13881.1		gb AAD34056.1 AF1	51819_1	emb CAB53709.1		
		similar to Human Sp2	protein (M97190)	Homo sapiens	(AF106473) leucine-	rich-domain inter-	acting protein 1; LeR 1	CTP synthetase	homolog [Mus	musculus]	D-E-A-D box protein	[Drosophila	melanogaster]	(AF213393) ATP-	binding cassette protein	[Mus musculus]	coded for by C. elegans	cDNA yk34b1.5; coded	for by C. elegans 1 1	coded for by C. elegans	cDNA yk46e8.3; coded	fo	adenylate kinase 2B	[Homo sapiens]	(AF151819) CGI-61	protein [Homo sapiens]	(AL110271)	hypothetical protein	Homo sapiens
		blastx.2			blastx.2		-	blastx			blastx.2			blastx.2		:	blastx.2						blastx.2		blastx.2		blastx.2		
		704			705			902			707			708			709						710		711		712		
		1044577			1044618			1044635			1044711			1044741			1044760						1044762		1044769		1044821		
		HSQEK12			HPDRZ16			HVCBC44			HPIAC22			HSODA53			HE9ML74						HPDRB76		HSYDISS		HOGCI31		

HNOKX86	1044893	713	blastx.2	(AF092138) HSPC033 [Homo sapiens]	gb AAD40380.1	100%	303	575
HTPHG81	1045117	714	blastx.2	(AL117402) hypothetical protein	emb[CAB55903.1]	100%	537	1538
HPCTV49	1045379	715	blastx.2	ribosomal protein	gb AAA36589.1	100%	127	444
нндрк68	1045464	716	blastx.2	(AL020993) dJ506.1 (casein kinase 1,	emb CAA15888.1	89% 100%	124	858
				epsilon) [Homo sapiens]		·		
HNOKM3 8	1045500	717	blastx.2	transducin (beta) like 1 protein [Homo sapiens]	emb CAA73319.1	%66	120	1640
HHFUN47	1045842	718	blastx.2	(AF099028) putative transmembrane protein cmp44E [Drosophila melanogaster]	gb AAD12254.1	49% 52% 27%	1007	1135 952 2368
HE9EJ82	1046856	722	blastx.2	beta-signal sequence receptor [Homo sapiens]	dbj BAA07206.1	%96	24	173
норнх68	1047137	724	blastx.2	(AK000826) unnamed protein product [Homo	dbj BAA91390.1	%66	528	1220

	814	786	586	786	802	404	200	935	687	554			6	90	,	1004	V 4 7	///6	1119	1432	401		3,0	7463		,,,	445		
	335	124	206	586	770	2 5	101	762	589	114			1	16/	18	2, 5	706	939	1081	509	141		1	40/		;	3		
	100%	100%	100%	94%	7007	2/9/	%07	38%	30%	%55				%86		%66	%96	46%	46%	36%	30%	•		100%			100%		
	gb AAA18537.1	gb AAF00499.1 AF1 82844 1	ahi 4 B B 69312 11	golden in the second se				-		emb CAB04731.1				emb CAA31134.1		dbj BAA92312.1				emb CAB11047.1				emb CAB75652.1			gb AAD38498.1 AF1	16272_1	
saniens	omo sapiens]	(AF182844) VPS28	+	(Artuil/92) cell cycle	progression 2 protein	[Homo sapiens]				predicted using	Genefinder	[Caenorhabditis	elegans]	ORF X (AA 1 - 393)	Escherichia coli	(AB034912) WD-	repeat like sequence	[Homo sapiens]	family array	putative RNA-binding	protein	[Schizosaccharomyces	pombe]	(AL157427)	hypothetical protein	[Homo sapiens]	(AF116272) T-cell	activation protein	Homo sapiens
	blastx.2	blastx.2	0.11.	DIASTX.2						blastx.2	-			blastx.2		blastx.2				blastx.2		-		blastx.2			blastx.2		
	725	726	200	17/						728				729		730				731	<u> </u>			732			733		
	1047169	1047212	, , , , ,	1047381						1047403				1047473		1047483				1047634	10011			1047646			1047663		
	HOPKTSO	HMCFK45	\dagger	HPIAN63						HNSME49			***	HWEAC64		HOCOI51	222			TOBVETS	CICALION			HMAEL73			HNOKE42		

			, 	T			· · · · ·	
181 221	305	399	958 249 907	729	1959	1279	1231	934
180	69	166	233 52 503	169	778	446	17	191
93%	100%	.87%	99% 91% 32%	94%	100%	30%	46%	100%
pir A00119 CCBO11	dbj BAA83718.1	gb AAC64583.1	dbj BAA13448.1	sp G545790 G545790	gb AAA98961.1	emb CAB11599.1	dbj BAA91513.1	gb AAB81515.1
ubiquinolcytochrome- c reductase (EC 1.10.2.2) 11K protein - bovine	(AB016092) RNA binding protein [Homo sapiens]	(AF091242) ATP sulfurylase/APS kinase 2 [Homo sapiens]	Similar to Human C219-reactive peptide (L34688) [Homo sapiens]	DARPP- 32=DOPAMINE AND CAMP-REGULATED PHOSPHOPROTEIN.	transketolase [Homo sapiens]	hypothetical protein [Schizosaccharomyces pombe]	(AK001123) unnamed protein product [Homo sapiens]	(AF022815) proteasome subunit XAPC7 [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
734	736	737	738	739	740	741	742	743
1047670	1047820	1047848	1047937	1048009	1048188	1048300	1048427	1048595
HOFAE31	HNOAC93	HMEJA45	HLDASII	HWMJB31	HAZAA31	HOCMC83	HCFCS40	HVCAA65

HSPSI76	1048635	744	blastx.2	ylase it	gb AAB71339.1	%66	807	0/61
HVVCH35	1048658	745	blastx.2	(AF177385) cytochrome c oxidase	gb AAF05313.1 AF1 77385_1	%66	436	1233
				assembly protein isoform 2 [Homo				
HOFNI66	1048739	746	blastx.2	sapiens signal peptidase complex 25 kDa	gb AAA21254.1	%56	33	710
				subunit [Canis			0,00	2277
HMSKI90	1048792	747	blastx.2	unknown protein [Homo sapiens]	gb AAA88038.1	39% 28%	1452	1180
	<u> </u>			-		100%	327	310
HSYBI49	1049151	748	blastx.2	(AF101051)	gb AAD16433.1	100%	173	802
				senescence-associated epithelial membrane				
HTXSN37	1049372	749	blastx.2	(AL080159) hypothetical protein	emb CAB45750.1	47%	1154	1 86 1 530
HMUBT31	1049466	750	blastx.2	Homo sapiensi serine palmitoyltransferase,	emb CAA69941.1	100%	11	1429
				subunit I [Homo sapiens]		/800,	713	611
HFPDO90	1049644	751	blastx.2	Similarity to yeast hypothetical protein	emb CAA94801.1		010	
	:			PIR accession number				

		1139	_			-	1186			2415					4	1273	7		810			<u></u>			1300	416
		213 1067	11				212	 .		643						161	1818	405	121	203	465	3017	1464	2939	371	m
		%96 %96	100%				%001			%66						100%	%66	97%	53%	31%	32%	40%	28%	46%	%86	%96
		emb CAA00829.1	emb CAB42866.1				gb AAA36043.1			splQ16236 NFL2_HU MAN						gb[AAB58075.1]	dbj BAA08565.1				-				gb AAC39776.1	-
1 1 this gene; cDNA	EST yk504c7.3 comes from this gene; cDN	urokinase [synthetic construct]	(AL031733) dJ455J7.1	(cellular repressor of	E1A-stimulated genes	CREG) [Homo sapiens]	interferon regulatory	factor 1 [Homo	sapiens]	NUCLEAR FACTOR ERYTHROID 2	RELATED FACTOR 2	(NF-E2 1	ERYTHROID	DERIVED 2, LIKE 2)	(HEBPI).	ESX [Homo sapiens]	DNA binding protein	[Homo sapiens]						•	(AF016903) agrin	precursor [Homo
		blastx.2	blastx.2				blastx.2			blastx.2						blastx.2	blastx.2	•							blastx.2	
		752	753			-	754			755						756	757								758	
		1050102	1050167				1050256			1050282						1050419	1050536								1050553	i
		HVVCB79	HDPGR19				HVVDX63			HNOJA87						HTPGI65	HHFGN14								HTAIN76	

1252	1291	416	440	721	557	430	999				648	196	557	597			574 672	695		298		403	:	522		
614	614	n	m	386	426	86	465	19	1125	m	541	155	42	961			188 571	174		245		99		244	53	385
36%	32%	45%	31%	38%	40%	76%	32%	35%	27%	28%	40%	28%	100%	%56			91% 45%	%58		%66		100%		82%	%96	23%
							-						emb CAA51467.1	gb AAB17261.1			gb AAC78495.1	gb AAF36134.1 AF1	51048 1	dbj BAA05167.1		gb AAB84248.1		emb CAA57478.1		
sapiens	i			•									caltractin [Homo sapiens]	serine/threonine protein	kinase Krs-1 [Homo	sapiens	annexin I [Oryctolagus cuniculus]	(AF151048) HSPC214	Homo sapiens	V-1 protein [Rattus	norvegicus	(AF028823) Tax interaction protein 1	[Homo sapiens]	p0071 protein [Homo	sapiens]	
			•										blastx.2	blastx.2		,	blastx.2	blastx.2		blastx.2	•	blastx.2		blastx.2		
		_											759	760			761	762		292		764		765		
													1050702	1050767			1050969	1051095		1051115		1051246		1051256		
													HLYBO89	HMEKJ82			HOFNX17	HAOSY21		HMWDB3	9.	HAOSZ53		HSODP14		

452 580	187 300	-		098 988	18 527		162 676	175 360			365 658	94 1377	5 82	3194 3250		2905 2952		184 2094	93 668	18 968	20 583
	34%			8 8	100%	37% 2	100% 3					100%	65%		40% 6		42% 6	100%	100%	%96	100%
		-	dbj BAA91205.1		dbj BAA90946.1		gb AAB60490.1		****			emb CAB63856.1	emb CAA92991.1					emb CAA24748.1	gb AAD34108.1 AF1 51871 1	gb AAA59925.1	emb CAB43305.1
			(AK000496) unnamed	protein product [Homo sapiens]	(AK000101) unnamed	protein product [Homo sapiens]	cellular nucleic acid	binding protein [Mus	musculus]		-	(AJ251914) putative RNA helicase [Sus scrofa]	Huntington Disease	(HD) gene exon 1	[Homo sapiens]	•	•	mannitol permease [Escherichia coli]	(AF151871) CGI-113 protein [Homo sapiens]	neurofibromin [Homo	(AL050169) hypothetical protein
			blastx.2		blastx.2		blastx.2					blastx.2	blastx.2					blastx.2	blastx.2	blastx.2	blastx.2
			992		167		89/					692	770			•		771	772	773	774
			1051319	•	1051410		1051437					1051533	1051883			·	•	1051903	1051953	1051983	1052158
			HWLXZ72		HUSGQ45	,	HLDRA54			•		HNOJR48	HKIXH35					ннЕВ 192	HPTGB84	HVCCD05	HPYSC40

	8 844	59 652	6 852	50 745	2 1033 3 650		2 431		2 622		2 697	9 1289	
	158		226		632	_	282				512	609	
	100%	100%	100%	%06	52% 49%	40%	64%	····	87%		100%	100%	
	gb AAF14858.1 AF1 10775_1	gb AAA60588.1	dbj BAA22984.1	gb AAC50895.1	gb AAC36017.1		gb AAA92286.1		gb AAC39930.1		emb CAA34385.1	dbj BAA91907.1	
[Homo sapiens]	(AF110775) adrenal gland protein AD-002 Homo sapiens	sorcin CP-22 [Homo sapiens]	(AB000712) CPE- receptor [Homo sapiens]	CUG-BP/hNab50 [Homo sapiens]	(AC002397) C9 [Mus	[mmamm	nuclear protein essential for dosage	compensation [Caenorhabditis	(AF064801) multiple membrane spanning	receptor TRC8 [Homo sapiens]	54k protein (AA 1-504) [Canis familiaris]	(AK001782) unnamed protein product [Homo sapiens]	
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2		blastx.2		blastx.2		blastx.2	blastx.2	
	77.5	2776	777	778	780		781		782		783	784	
	1052261	1052553	1052557	1052593	1052874		1053037		1053164		1053171	1053173	
	HDDMT56	HVVAW0	HOGCS42	HAZAA59	HMALJ21		HLYAR61		HLTCQ80		HDTDU67	HMABL01	

2631 1813 889	653 518 521	610	532	220 587 431	869	808
1804 1301 815	39 42 42	95	374	77 486 387	150	164
99% 100% 60%	66% 28% 27%	100%	%06	100% 48% 53%	100%	100%
dbj BAA24179.2	gb AAB59189.1	dbj BAA11212.1	dbj BAA20422.1	emb CAA72272.1	gb AAA79920.1	gb AAA91892.1
(AB002405) LAK-4p [Homo sapiens]	pumilio protein [Drosophila melanogaster]	CIRP [Homo sapiens]	heparan sulfate 2- sulfotransferase [Cricetulus longicaudatus]	phosphoenolpyruvate carboxykinase (GTP) [Homo sapiens]	microfibril-associated glycoprotein [Homo sapiens]	casein kinase-II beta [Oryctolagus cuniculus]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
785	786	787	788	789	790	791
1053236	1053369	1053547	1053548	1053585	1053725	1053746
HACMU05 1053236	HOVDF79	HVVBJ54	HOFMUSO	HE9MO38	HOPJF55	НАОТЪ13

27 - +57 - 2004	1053973	793	blastx.2	p0071 protein [Homo	emb[CAA5/4/0.1]	9,0%		
-				sapiens	AN A C 23797 11	%66	178	1614
HFKHC64 10	1054015	794	blastx.2	(AF058448) herpesvirus entry	li			
				protein B [Homo				
				sapiens]	αh Δ Δ Δ 16256.11	%66	7	1063
HOPKO37 10	1054085	795	blastx.2	drebrin E. [Homo	المصميحيامة			
-1				Saplens	obl A AC17708.1	%68	90	2332
HNBV053 10	1054122	796	blastx.2	ferrordes orrord not d		46%	639	1403
	-				•	31%	1871	2380
						36%	883	1119
						%06	36	89
						31%	754	1101
						37%	952	1173
<u>нсоро</u> 11	1054196	798	blastx.2	cytochrome c oxidase subunit 3 [Homo	dbj BAA77671.1	%86	25	777
				sapiens	1 10000	1000%	226	873
HTFML39 1	1054230	799	blastx.2	secreted cyclophilin- like protein [Homo	gb AAA36601.1	80001	077	5
HOGCR32	1054235	008	blastx.2	similar to mouse CC1.	dbj BAA13194.1	%06	15	110
				[Homo sapiens]		35%	128	169
HECDI 60	1054288	801	blastx.2	(AF005038) secretory	gb AAB62723.2	%66	152	919
				carrier membrane		0000	7	2
HODEC13 1	1054400	802	blastx.2	(AC007059) Human	gb AAD19818.1	81%	m	722
				homolog of Mus wizi, protein				

	285	1989	1620	948	1985	1348	1682	979	1276	895	1333	1324	1273	895	856	1003	1147	979	484	1445	1676	1676	829	1769
	٢	1084	841	100	1908	2	1290	251	62	197	11	44	62	146	2	605	278	35	221	1308	1290	1437	969	1287
	97%	73%	45%	30%	25%	%86	%06	31%	27%	30%	24%	27%	79%	30%	79%	32%	24%	20%	34%	82%	76%	78%	78%	25%
	dbj BAA02807.1	gb AAC00205.1				dbj BAA22956.1																		
sapiens]	protein related N- ternimus of tre oncogene [Homo sapiens]	PRAJA1 [Mus	musculusi			(AB006625) The	himan homolog of a	monse imprinted gene.	Peg3. [Homo sapiens]				•											
	blastx.2	blastx.2				blastx.2																		
	813	814				815	3								_									
	1055174	1055248				1055304																		
	нтенР29	HNO1132				HEPHES?	70 117 1 177												-					· ·

1676	1499	1676	979	1499	1679	1676	1502	625	829	1048	1438	581	1191	1006		1310	1504	1518		303	770	689
1308	1305	1398	905	1425	1596	1608	1344	314	752	86	7	111	571	839	-	522	1433	154		33	3	. 201
792	32%	27%	36%	36%	39%	34%	28%	100%	43%	100%	100%	%26	28%	25%		42%	32%	%66		7000	0/66	%86
					-			dbj BAA91239.1		emb CAA55343.1	emb CAA40918.1	gb AAC51293.1				gb[AAC37427.1]	-	emb CAB61393.1		amble A B 6 5 1 10 11	اتختتوصحمامته	gb AAD45242.1 AF1
						•		(AK000538) unnamed	protein product [Homo sapiens]	EF-hand protein [Homo sapiens]	stromelysin-3 precursor [Homo sapiens]	Hlark [Homo sapiens]				myosin IC	[Dictyostelium discoideum]	(AL133070)	hypothetical protein	(A 1250865) TEGG 2	[Homo sapiens]	(AF118394) putative
								blastx.2		blastx.2	blastx.2	blastx.2				blastx.2	٠	blastx.2		bloch 2	Oldsich.2	blastx.2
								816	-	817	818	820				821	_	822		873	020	824
							,	1055381		1055426	1055439	1055467				1055480		1055582		1055632	700001	1055767
								HBNMF62	•	нтрно72	HOPKL18	HTXKD84				HTXRR82		HAIBU62		HDTA V20	(7141711	нсоор79

210	550	562				67.0	/661			650	823							168	777	440	1001	1001		3	766			
9/	53	263				ľ				Ť	7					-			6	<u></u>	100	188		ľ]
100%	2007	36%									45%							 %86	, 600	100%	,000	%00I			%86			
18394_1	1104 40470 11	emb[CAA94773.1]					gb AAD49722.1 AF1	0/100_1			emb CAA99881.1		-				•	emb CAA66186.1		gb AAC51319.1		gb[AAB51324.1]			gb AAD53962.1 AF1	8140/_1		
\vdash	protein [Homo sapiens]	Similarity to EGF	FMBL: T02406 comes	from this gene	Caenorhabditis	elegans	(AF167160) protein	inhibitor of activated	STAT-1 [Homo	sapiens	similar to ankyrin	motifs; cDNA EST	CEMSH89F comes	from this 1 1 cDNA	EST EMBL:D33056	comes from this gene;	cDNA EST E	fsh-like protein [Mus	musculus	Wnt7a protein [Homo	sapiens]	B-cell receptor	associated protein	[Homo sapiens]	(AF181467) protein Z-	dependent protease	inhibitor precursor	[Homo sapiens]
		blastx.2					blastx.2				blastx.2		•					blastx.2		blastx.2		blastx.2			blastx.2			
		825					826				827							828		829		830			831			
		1055899					1056000				1056097		٠				•	1056102		1056104		1056275			1056290			
		HAMHIN1	2	_			19XIGOH				HNTRS57							HUKFL74		HTEBF05		HOPKD19			HLICR58			

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111	1436	327		3060	2118	2133	2166	2166	2118	2118	1632	2118	2109	2117	940	934	931	4139	1251	4190	3694	771	841	1587	3154	771	4119
7	366	265		1036	1036	1033	1036	1264	1429	1486	1036	1594	1327	1032	743	743	899	3777	1015	3672	3572	658	728	1432	2846	565	4021
100%	100%	33%		%56	42%	43%	45%	46%	44%	45%	44%	38%	36%	792	38%	35%	34%	37%	31%	72%	36%	44%	44%	29%	73%	32%	40%
sp Q9Y6Y5 Q9Y6Y5	gb AAC98506.1			gb AAC39658.1				•								÷											
IDN4-GGTR14 PROTEIN.	(AF095448) putative G	protein-coupled	receptor [Homo sapiens]	(AF018081) type XVIII	collagen [Homo	sapiens	•	-														-					
blastx.2	blastx.2			blastx.2																							
833	834			835																							
1056400	1056407			1056454		٠.																					
HAJAH48 1056400	HSKJC61			HEEBK29			•																				

784	901	910	2905	3607	3161	3161	3971	3161	3731	3164	1140	1104	1056	1140	1053	1140	1140	1056	1164	1050	1230	1368	1566	1566	1692	1800	1632	1859	1608
229	749	707	2753	3524	3120	3087	3789	3108	3579	3087	2129	2108	2129	2039	2129	2015	2105	2129	2108	1910	2111	2108	2111	2111	2039	2105	2108	2707	2108
38%	37%	27%	35%	32%	20%	40%	27%	20%	33%	40%	40%	37%	35%	38%	37%	37%	36%	35%	32%	35%	38%	37%	38%	37%	43%	43%	38%	30%	37%
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2117	2680	1468	2707	4790	2997	3599	2394	3629	4135	2394	2457	3692	2020	3997	3015	3877	3862	3163	2988	4843	2394	1183	894	2825	178	2892	2157	4128	4131
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2801 2	10	1	. 271			1,1	<u>م</u>	1	4°C		163		/009						290					7	26	79	1031
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	emb CAA58449.1		emb CAB55680.1				gb AAC69883.1		pir A60345 A60345		11 + 11 + 00000 CE	gb AAF08220.1 AF1	51793_1		pir A32891 A32891									dbj BAA22621.1	ob AA58460.1	-	
	Lutheran blood group		8) dJ479J7.1	(similar to	CHONDROMODULI	N-1) [Homo sapiens]	(AF094760) RFXANK	[Homo sapiens]	protein-tyrosine-	phosphatase (EC	3.1.3.48) IIA - numan	(AF151793) ALG-2	interacting protein 1	[Homo sapiens]	finger protein 1.	alocental himan	piacentai - manas							GATA-6 [Homo	sapicins protein disulfide	isomerase-related	protein [Homo sapiens]
	blastx.2		blastx.2				blastx.2		blastx.2			blastx.2		,	blacty 2	- W. C.								blastx.2	110000	Oldstv.2	
	836		837				838		839			840	?		9.4.1	7 -								842	5,5	0 1 0	
	1056617		1056625			•	1056654		1056666			1056671			1055570	7/00001								1056736	7,2,0	1020/04	
	HCHWM1	,	HE8NG02				HWHKD2	2	HOCPJ87			UAAAA50	Constant		01100011	HOSBITS								HUSGX12		HPDRG02	

1327	1660	1316	650	983	831	205	1833 1875 1779 1809 1782 1812 1812
1055	239	111	87	702	4	300	727 556 727 703 703 706 892
40%	100%	%86	100%	32%	%86	75% 62%	30% 26% 26% 24% 24% 24% 24% 26%
	dbj BAA78677.1	gb AAD32671.1 AF1 40242_1	gb AAA91463.1	gb AAB61297.1	emb CAA31282.1	dbj BAA85438.1	emb CAA62188.1
	(AB022663) HFB30 [Homo sapiens]	(AF140242) encephalopsin [Homo sapiens]	VEGF related factor isoform VRF167 precursor [Homo sapiens]	(AF003944) ovalbumin upstream promoter beta nuclear receptor rCOUPb [Rattus norvegicus]	ear-2 gene product [Homo sapiens]	(AP000616) similar to RING-H2 finger protein RHA1a (AF078683) [Oryza sativa]	Protein sequence and annotation available soon via Swiss-Prot; 1 [Homo sapiens]
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	844	845	846	847	848	849	850
	1056767	1056774	1056786	1056801	1056804	1056810	1056839
	HKZBB48	HHAWB1 9	HHATP38	HVVBE07	нканв85	HUSXA15	HNOCH54

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727	724	733	883	856	28	883	865	856	886	730	4	1	715	7	145	31	946		31	16	382	745	22	-	7		424	10	724
24%	21%	24%	25%	76%	22%	23%	21%	22%	23%	22%	24%	25%	22%	32%	31%	24%	23%	28%	25%	22%	23%	23%	70%	792	28%	26%	24%	30%	23%
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259	22	430	427	1054	250	427	430		7	427	352	730	250	424	427	1108	415	430	313	427	322	322	430	343	1648	430	412	430	1369
25%	31%	25%	34%	21%	35%	24%	24%	30%	25%	23%	78%	23%	31%	34%	35%	24%	32%	79%	76%	73%	27%	27%	30%	30%	36%	33%	31%	22%	22%
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1273	424	412	961	427	430	409	634	430	1123	427	430	1648	1267	427	421	418	1651	1648	1072	1288	388	424	499	1621	1375	1279	427	430	430
25%	30%	28%	25%	30%	22%	22%	25%	21%	21%	31%	31%	31%	72%	27%	78%	79%	78%	30%	22%	24%	73%	72%	25%	79%	22%	30%	30%	32%	25%
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1660	1648	1648	1333	415	1648	1273	1006	1009	418	1282	1582	430	148	1117	1648	1123	16	1618	1123	1012	1390	382	1369	352	427	427	430	427	430
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424	1648	1648	1018	1651	1648	1384	1648	1648	1012	430	1669	1393	1648	1123	1369	1648	427	1648	1057	961	1357	1648	926	1222	724	4	1390	1123	1648
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206	1288	1075	1120	1207	409	28	1285	16	1648	1648	1463	1072	1369	1648	1642	1012	338	1651	1174	1357	1369	1375	754	440	427	-	124	1651	1648
70%	25%	70%	22%	21%	76%	25%	22%	79%	72%	73%	37%	18%	21%	24%	25%	25%	27%	75%	23%	79%	23%	21%	16%	79%	24%	25%	25%	23%	. 22%
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2095	144	1314	1362	144	1368	1827	1091	1458	1929	1827	533	2018	1236	1175	1827	1827	1509	2092	1837	1692	528	1731	2060	1440	1554	1467	1830	2078	1321
2018	1	1195	979	67	1123	1648	897	1318	1468	1648	477	1980	1120	993	1651	1648	1375	1910	1733	1648	397	1369	1878	1336	1327	1120	1648	2031	1256
42%	25%	34%	75%	34%	30%	70%	78%	25%	22%	70%	47%	23%	23%	33%	22%	75%	24%	24%	72%	23%	73%	70%	27%	31%	25%	23%	21%	43%	40%
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849	1928	393	1236	650	848		480	229		469	343	319	313	274	403	313	313	321	403	155	131	155	176	155
814	1863	127	1120	36	90		26	5		=======================================	14	70	71	14	14	14	38	136	299	9	48	45	63	54
20%	45%	27%	25%	100%	100%		100%	%89		%66	42%	47%	20%	48%	34%	38%	38%	17%	77%	37%	42%	32%	45%	38%
				gb[AAA03246.1]	gb AAA87064.1		dbj BAA23363.1	dbj BAA91205.1		emb[CAB70782.1]	-							gb AAD31764.1 AF1	21963 1	i				
				insulin-like growth factor binding protein 2	Homo sapiens platelet-endothelial	tetraspan antigen 3 [Homo saniens]	OTK27 [Homo	(AK000496) unnamed	protein product [Homo sanjens]	(AI,137516)	hypothetical protein	[Homo saniens]	Litomo saprema					(AF121963) receptor	tyrosine kinase	precursor [Gallus	gallis]	Eurita		
				blastx.2	blastx.2		blastx.2	blastx.2		hlastx 2								hiacty 2	7:V367					
				851	852	}	853	854		855	3							856	<u>}</u>					
				1056856	1056862		1056875	1056927		1056000	0660001							1057000	600/601					-
				HOCPP16	HNOAH83	Correspond	HE20U10	HODGM4	9	TIONEDAA	HODER44							21777777	HCHINDIZ					•

424 759	400 815	2206	2084	1687	1690	1687	1705	1768	1681	1687	1687	886	1027	122/		1155	828	9/9					713	1274
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%06 86	39% 100%	%86	34%	36%	35%	37%	36%	36%	34%	34%	34%	38%	1000/	100%		%86	22%	77%					%89	47%
gb AAA.59545.1		gb AAA51973.1				-						gb AAC32982.1	11 0 0 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	emolCAB43289.1		16109B 16109B zid	,	PHA AF01349 11AC0	05003 1	i			emb CAA91418.1	
myelin-associated glycoprotein precursor	[Homo sapiens]	carboxyl ester lipase	[Homo sapiens]									junctional adhesion molecule [Mus	(AT 050143)	(ALUSU143)	hypothetical protein [Homo sapiens]	transcription regulatory	protein Evi-1, short	(AC005003) similar to	zinc finger protein	MAZ [Homo sapiens];	similar to AAB04121.1	(PID:g995935)	Homology with Squid	retinal-binding protein
blastx.2		blastx.2										blastx.2		blastx.2		blastx.2		blacty 2					blastx.2	
857		858	•									859	3	098	-	861		698	3				863	
1057157		1057170										1057212	0,000	1057219		1057260		1057272	7/7/001				1057307	
НВНАС29		HTPFW87				·						HISAF60	0000	HRABO80		HADDF30		HDA A 858	account of the				HSDZG15	

				(PIR Acc. No. 11	11 000000	2017	4	1196
HPDVY52	1057393	864	blastx.2	laminin alpha 5 chain	gb AAC33430.1	47%	1165	1380
HHESW02	1057478	865	blastx.2	(AF062075) leupaxin	gb AAC16014.1	100%	968	1252 1258
HVVBW8	1057561	998	blastx.2	(AB003184) ISLR	dbj BAA22848.1	100%	342	1625
1000	1057707	898	blastx 2	prolargin [Homo	gb AAC18782.1	32%	353	1228
HSLJC80	1611cor	900	Z.W.Corro	sanjens]		32%	868	1573
				Correction		28%	989	1420
						78%	1109	1657
						75%	1663	1980
-						34%	262	35/
HEOAD12	1057842	698	blastx.2	(AF116865) hedgehog-	gb AAD31172.1 AF1	%02	. 533	1069
				interacting protein	16865_1			
		,		Mus musculus	* **	70001	130	1272
HVVBT41	1057880	870	blastx.2	(AL035587) dJ475N16.1 (CTG4A)	emb CAB/>301.1	100%	r t	1
				Homo sapiens		,,,,,,	220	650
HAZAE42	1057915	871	blastx.2	cellular retinol binding protein [Homo sapiens]	emb CAA30318.1	100%	667	600
HPRBN60	1057948	872	blastx.2	(AF002282) alpha-	gb AAC16672.1	%66	72	1019
				LIM protein [Homo				
				sapiens		2000		7.80
HOVCZ68	1057958	873	blastx.2	follistatin 1 precursor -	pir A32141 A32141	%66	1	100
				numan	1.104 4.40105 11	7000	မ	1021
H6EDF71	1057979	874	blastx.2	antigenic surface determinant OA3	emo CAA49190.1	0/66	*	

				[Homo sapiens]				
HCOOF60	1058001	875	blastx.2	ubiquitin carrier protein E2 - human	pir B42856 B42856	%66	133	855
HVVBH81	1058059	928	blastx.2	hnRNP U protein [Homo sapiens]	emb CAA46472.1	%66	227	2644
HSPTJ51	1058172	877	blastx.2	SWI/SNF complex 155	gb AAC50693.1	%56	51	1004
				KDa subunit [Homo		%06	51	1043
				sapiens		29%	45	1055
HOPKC80	1058174	878	blastx.2	(AJ011497) Claudin-7 [Homo sapiens]	emb CAA09626.1	%66	348	086
HOCPT58	1058250	879	blastx.2	(AF037261) SH3-	gb AAC09244.1	100%	121	765
				containing adaptor		94%	19	126
				molecule-1 [Homo saniens]				
HPCTP25	1058287	088	blastx.2	gamma-	gb AAB18827.1	%86	72	1553
				aminobutyraldehyde				
	٠			dehydrogenase [Homo				
HODKWS	1058305	881	blocty 2	(A BOO8780) 22h7	41:11D 4 4 20050 11	2000	5	100
2			7:57:57:57	protein [Homo sapiens]	T'ACOATATAIN	76%	× n	666
HCONB89	1058316	882	blastx.2	acid ceramidase [Homo	gb AAC50907.1	%66	118	1302
				sapiens]				
HBPND88	1058432	884	blastx.2	carboxyl terminal LIM	gb[AAC05580.1]	%66	142	1128
				domain protein [Homo	•			
H6EEO05	1058438	885	blastx.2	(AL117423)	emb CAB55915.1	%66	100	1527
				hypothetical protein	-			
				LICHTO SAPIETIS				
HFIVK61	1058451	988	blastx.2	interferon beta 2a	emb CAA00839.1	%66	214	111

				[Homo sapiens]	٠			
HVVBX28	1058458	887	blastx.2	SH3 domain-containing	gb AAC50592.1	%96	53	493
				protein SH3P17 [Homo	-	46%	308	466
				sapiens	-	36%	104	301
HVVBI38	1058475	888	blastx.2	factor H [Homo	emb CAA68704.1	%66	294	2291
				sapiens	-	25%	336	2264
,				1	-	79%	294	2291
					-	72%	501	2219
						78%	1083	2288
						27%	1185	2099
HUKEJ46	1058539	688	blastx.2	(AF081507) signaling	gb AAC33967.1	%86	479	1165
	•			molecule LEFTY-B		% 66	70	489
				Homo sapiens		1,000	1	3
HDTFT47	1058588	890	blastx.2	lymphocyte antigen	gb AAA36236.1	%86	7	768
				[Homo sapiens]		91%	258	398
HSSFS71	1058596	891	blastx.2	ESP1/CRP2 [Homo	dbj BAA07703.1	%56	29	460
				sapiens		71%	41	265
				1		34%	219	512
НЅЪЈН63	1058612	892	blastx.2	(AB045180) toll-like	dbj BAB19259.1	95%	192	1358
				receptor 9 [Homo		100%	107	190
				sapiens		52%	128	184
	-			1		45%	122	181
						28%	131	181
HAHGD24	1058622	893	blastx.2	laminin alpha 5 chain [Mus musculus]	gb AAC53430.1	61%	08	1024
HTAEV85	1058723	894	blastx.2	(AJ005566) SPR2H	emb CAA06595.1	43%	408	256
				protein [Mus musculus]		64%	2500	2459
	•.					20%	1784	1719
	-				•	35%	544	485
HPMME44	1058928	895	blastx.2	HCMVUL126 [human	emb CAA35328.1	%96	78	329
	4							

		L	-	herpesvirus 5				
HSPSG28	1058977	897	blastx.2	26S protease subunit [Sus scrofa]	emb CAA61863.1	%88 %66	483	1679
			, 					
HTGFW12	1059006	868	blastx.2	homolog of yeast mutL	gb AAA63923.1	%86	83	1438
		·		gene [Homo sapiens]		94%	1522	2670
HOFMT75	1059050	668	blastx.2	cathepsin D [Homo	gb AAA51922.1	87%	7	456
יייייייייייייייייייייייייייייייייייייי	20000	300		sapiens	-	%08	414	743
HAKINB1/	1059085	906	blastx.2	HCMVUL126 [human herpesvirus 5]	emb CAA35328.1	%86	241	492
HPDRG65	1059102	10	blastx.2	(AK000496) unnamed	dbj BAA91205.1	%69	434	273
				protein product [Homo		%69	253	128
		1		sapiens		83%	288	253
HAOSG15	1059145	905	blastx.2	(AF026291)	gb AAC96010.1	100%	160	1776
				chaperonin containing				
				t-complex polypeptide		 -		
				delta [Homo sapiens]		,		
HHEND31	1059180	903	blastx.2	(AF117330) unknown	gb AAD26207.1 AF1	73%	175	1827
				[Rattus norvegicus]	17330 1			
HBMSN62	1059186	904	blastx.2	ZZ:beta-Gal' IgG-	gb AAB00807.1	%98	1102	1251
	:			binding fusion protein	•			
				[unidentified cloning 1			,	_
HOVCJ46	1059241	905	blastx.2	(AF037272) WAP	gb AAC40055.1	87%	41	214
				rour-disultide core		70%	175	336

	1348	1191	1698		685	1996	208	2310	955	}	1048		496
	92	436	427	1	2	212	188	2209	158	901	125		275
	%66	100%	100%		100%	100%	37%	32%	7000	97.66	%16	-	52%
	gb AAD34341.1 AF1 17646_1	gb AAC02781.1	11 120001 1 121	db] BAA13404.1	gb AAA50779.1	1-14 4 4 2 5 0 6 2 1 1	go AAA33303.1			gb[AAC39802.1]	gb AAF34411.1		emb CAA20980.2
domain protein [Rattus	long CBL-	omoH]		Similar to Human tesican (S33293)	phosphatase [Homo	sapiens	hematopoietic cell	pnopnatase [1101110 sapiens]		(AF062317) p120 catenin isoform 1B	(AF169284) LIM and	cysteine-rich domains	(AL031629) similar to RNA recognition motif.
	blastx.2	blastx.2		blastx.2	blastx.2		blastx.2			blastx.2	blastx.2		blastx.2
	906	806		606	910		911			912	913	;	914
	1059394	1059532		1059626	1059692		1059710			1059743	1050764	10000	1059784
	HWLEP14	HDABR73		HOGAJ24	HTJMV05		HOOIL70			HUNAE87	1001I	HIFHOUI	HDPWK69

1267	1216	1213	832	635	1150 1380	959.	581	808	189 308
65	575	215	23	447	1362 1487	18	207	197	189
%66	%86	100%	62%	100%	43% 47%	91%	39%	%62	70%
gb AAC12708.1	emb CAB56027.1	gb AAD30062.1	gb AAD12719.1	gb AAA58350.1	gb AAA88036.1	emb CAB70887.1	gb AAB41727.1	dbj BAA07295.1	emb CAA55356.1
(AF008551) aurora- related kinase 1 [Homo sapiens]	(AL117639) hypothetical protein [Homo sapiens]	(AF132856) suppressor of G2 allele of skp1 homolog [Homo sapiens]	(AF078798) extracellular signal- regulated kinase 7; ERK7 [Rattus	replication protein A 14kDa subunit [Homo sapiens]	unknown protein [Homo sapiens]	(AL137714) hypothetical protein [Homo sapiens]	PF20 [Chlamydomonas reinhardtii]	ATPase subunit 6 [Homo sapiens]	sialidase [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
915	917	918	919	920	921	922	923	924	925
1059849	1059967	1059969	1060137	1060193	1060382	1060391	1060415	1060495	1060656
нРСОК03	HOOJC15	норг647	HBIPC05	НОРЛЗ2	HNOBN20	HOCPX74	HTEPP27	HLYAJ79	HC00Q11

blastx.2
blastx.2 GTP binding protein
[Mus musculus]
blastx.2 hypothetical protein 2
(rRNA external
FRE-7 Homo sapiens
blastx.2 26S PROTEASOME
SUBUNIT S2 (P97)
(TUMOR 1 1
blastx.2 MRAS2 gene product
blastx.2 matrin 3 [Rattus
norvegicus
•
•
\dagger
blastx.2 (AJ000414) Cuc42=
interacting protein 4
\dagger
blastx.2 (ALV2212

				dJ1119A7.1				
				(mitochondrial		-		
				thioredoxin) [Homo	-			
			·	sapiens				
HWLLG38	1061388	936	blastx.2	neurocalcin [Bos	dbj BAA01706.1	100%	114	692
				taurus]				
HTPCH84	1061466	937	blastx.2	(AF104419) decoy	gb AAD03056.1	%66	103	945
	_			receptor 3 [Homo		100%	938	1000
				sapiens]				
HLGDA34	1061543	939	blastx.2	(AK000496) unnamed	dbj BAA91205.1	71%	3	254
		_		protein product [Homo		73%	1766	1641
				sapiens				
HAICB08	1061629	940	blastx.2	rhoHP1 [Homo	dbj BAA19652.1	100%	3	326
				sapiens				
HVVAW6	1061694	941	blastx.2	(AF151882) CGI-124	gb AAD34119.1 AF1	100%	244	741
0				protein [Homo sapiens]	51882_1			-
HPDVB50	1061708	942	blastx.2	(AJ002030)	emb CAA05152.1	100%	33	206
				progresterone binding				
_				protein [Homo sapiens]				
HAZAY40	1061765	943	blastx.2	tight junction (zonula	gb AAA02891.1	100%	3	1631
				occludens) protein ZO-				
7				1 Homo sapiens				
HACMZ51	1061766	944	blastx.2	contains 10 ankyrin-	gb AAC96986.1	33%	118	957
				like repeats; similar to		78%	73	726
	•			human ankyrin, 1		32%	100	657
				bursaria Chlorella virus		30%	88	561
				11				
HUXAL63 1061790	1061790	945	blastx.2	100 kDa protein	emb CAA45756.1	%86	2	1300
				rations Horvegicus				

953	941	398	398	941	914	941	676	941	914	686	929	676	941	167	929	149	149	377	155	914	149	365	155	143	371	944	143	188	215
m	447	27	21	969	693	969	681	999	693	999	684	657	999	12	969	9	15	6	m	829	3	6	c	m	246	837	m	15	66
%66	37%	32%	34%	45%	40%	41%	42%	38%	36%	32%	34%	34%	34%	43%	35%	45%	47%	31%	46%	35%	40%	73%	35%	38%	40%	41%	34%	29%	78%
gb[AAB37459.1]					-					-			-1-2-				-			•									
latent transforming	growth factor-beta-	binding protein-2, 1 1	(,	•																						
hlastx 2			,											•		•							-						
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252	468	1130	1290	1323	1528	1805	2170	1553	2338		1234	1165			1654			1535		
280	427	519	181	1141	1460	1710	1883	1506	284		200	95			209		·	862		
35%	27%	%66	%66 %98	27%	34%	33%	27%	20%	%66		%66	47%			100%			%16		-
		gb AAA30514.1	dbj BAA91446.1	,					gb AAC14573.1		gb AAD46623.2 AF1 62680 1	gb AAF21308.1			gb AAA84934.1			emb CAB43253.1		
		factor activating exoenzyme S [Bos taurus]	(AK000963) unnamed	sapiens]					(AF059617) serum-	inducible kinase [Homo sapiens]	(AF162680) STRIN protein [Homo sapiens]	(AF113596) mosaic	serine protease epitheliasin [Mus	musculus	retinoic acid- and	interferon-inducible	Homo sapiens	(AL050060)	hypothetical protein	Licomo sapiens]
		blastx.2	blastx.2						blastx.2	:	blastx.2	blastx.2			blastx.2			blastx.2		-
		947	948						946		950	951			952			954		
		1061935	1062057						1062079		1062084	1062123			1062139			1062309		
		HMALF63	HACMR36			-			HAOSM08	,	HTEFM89	HCHAK72			HPAME12	-		HCE4F10		

	790 251 96 63	248 887	1419 571	507	443	2129	2044	1663	1297	1070	10/9	7561		559	1
	191 87 25 1	361 1093	574 416	1/1	7/7	417	1997	c	188	9 5	/971	156		203	7
	100% 69% 87% 57%	97% 24%	99%	%/%	%66	38%	47%	7000	30%	%97	100%	%66		85% 91%	82%
	gb AAC39582.1	emb CAA51320.1	gb[AAC28019.1		dbj BAA21881.1	oblA AD05305.11			gb AAD43756.1 AF1 52495_1		gb AAF24048.1 AF0 90934_1	gb AAA36528.1		emb CAA73944.1	
- :	(AF007170) unknown [Homo sapiens]	endonuclease G [Bos	(AF062006) orphan G protein-coupled	receptor HG38 [Homo sapiens]	(AB006679) ATP binding protein [Homo	Sapiens (ATT11060) lotmonhilin	2 splice variant basae	Bos taurus	(AF152495) protocadherin beta 2	Homo sapiens	(AF090934) PRO0518 [Homo sapiens]	protein tyrosine phosphatase (EC	3.1.3.48) [Homo	latent TGF-beta binding protein-4	[Homo sapiens]
	blastx.2	blastx.2	blastx.2		blastx.2		blastx.2		blastx.2		blastx.2	blastx.2		blastx.2	
	955	956	957		958	1	959		961		962.	963		964	
	1062328	1062346	1062369		1062431		1062435		1062544		1062574	1062586		1062626	
	HBCBE63	HSSJ019	HE8NQ23		HTELJ95		HTPCP50		HFKIT82		HE2LW42	HOGCE44		HOVJJ72	

1728	363	1296	885	1869	1920	276	1926	198	1761	1734	1761	1881	198	1116	1743	198	153	583	268	1164	268	298	268	925	280	268	328	301	493
562		16	16	1537	1132	4	1465	H	1507	1435	1474	1366	4	553	1453	34	16	212	209	562	215	215	206	410	215	215	212	212	245
30%	45%	29%	30%	33%	33%	41%	32%	48%	38%	35%	36%	73%	47%	73%	38%	36%	34%	23%	%09	25%	%99	23%	42%	32%	. 52%	25%	38%	40%	29%
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346	307	271	210	265	310	921	1930	1861	265	460	1838	1842	1891	1450	95	406	584	569	1421	1969	1421	938	1352	1493	1266	320	1392
215	212	209	4	215	212	841	1766	1667	230	215	1764	1792	1859	1391	12	254	549	838	1633	2007	1621	1024	1552	1525	1295	15	4
37%	37%	42%	29%	25%	30%	29%	32%	32%	28%	762	26%	28%	63%	40%	37%	37%	%99	87%	34%	%69	30%	38%	30%	93%	70%	%001	%56
																								-		gb AAA35914.1	emb CAA79635.1
								`																		guanine nucleotide exchange factor [Homo	sapiens thrombospondin-4
								•																		blastx.2	blastx.2
																										965	996
																										1062628	1062629
																										HODBT14	HBOEB83

	2 247	2 247	2 247	337	514	247	247	490	468	247	259												313		1112		814	
	7						_		250		_										_	250		253	171		188	
	74%	74%	73%	26%	40%	%69	%69	41%	78%	%29	%99	71%	73%	77%	71%	75%	70%	%69	71%	64%	63%	72%	38%	49%	%0\$		100%	
	gb AAA79179.1	· ·				-				,				-								-			dbj BAA91301.1		gb AAA58659.1	
[Homo sapiens]	repressor	transcriptional factor	[Homo sapiens]	1								-	-												(AK000642) unnamed	protein product [Homo sapiens]	high mobility group 2	•
	blastx.2																								blastx.2		blastx.2	
-	296			-																					896		696	
	1062631			•																					1062655		1062679	
	HODCT96																							-	HTXJE60		HUSIQ62	

1562	 	649	1010	0161	1063	958	1055	949	460	705	705	705	999	200	707	70.5	2 5	705	705	705	705	705	
975	 	365	360	200	728	728	735	734	119					7 -	-	- F	4	r4	<u> </u>	-	10	1	
100% 93%		%16	7000	0%66	7072	36%	36%	38%	27%	%86	3/2/	76%	010/	0170	13%	2007	12%0	74%	%9/	74%	75%	74%	
gb AAF29140.1 AF1 61525_1		dbj BAA08312.1		gb[AAD51367.1 AF1 77203_1	11 00 00 4 4 00 11	emb CAA40203.1				gb AAA59469.1												-	
(AF161525) HSPC177 [Homo sapiens]		LIMK-2 [Homo	sapiens]	(AF177203) cerebral cell adhesion molecule	Homo sapiens	extensin [Volvox	carteri			zinc finger protein	[Homo sapiens]											-	
blastx.2		blastx.2		blastx.2		blastx.2				blastx.2													
971		972		973		974				975													
1062718		1062743	21.14001	1062785		1062795				1062840													
HKBAK29	 	9CA-JYNGIT	III IMPONED	HFKHF51		HUCPE28				HPWAH30	1												•

74% 173% 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	705	705	705	705	705	705	705	705	705	705	705	705	705	705	705	705	705	815	815	815	815	815	815	815	815	815	815	. 815	815	815
72% 72% 71% 71% 71% 71% 71% 71% 71% 71% 71% 71	F		-	_		-		_					_		_	-	163	702	723	723	702	723	702	723	702	723	702	702	726	723
	74%	72%	73%	72%	71%	72%	71%	71%	71%	71%	%02	71%	71%	71%	%19	63%	%65	52%	21%	51%	45%	48%	45%	45%	39%	45%	39%	36%	46%	45%
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815	200	815	815	815	815	809	815	815	815	210	5	3	250		170	0/1	911	419	245	
702	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	702	723	723	723	726	726	126	726	10	1	7	c	4	3,5	3	m	42	~	1
36%	, , ,	36%	41%	11%	41%	42%	43%	130%	43%	%56	7020	%c%	7000	07.60	,800	93%	100%	100%	10007	100%
							-			gb AAA19775.1		gb AAB59501.1	11 00000 1 1 211 22	dbj BAAU/290.1		gb AAA24905.1	gb[AAB65437.1]	gb AAA82926.1	11 000001 4 701 6	emb CAA52882.1
					•	_			•	cytochrome b [Homo	1	-beta protein	Homo sapiens	NADH dehydrogenase subunit 1 [Homo	sapiens	TraC protein (gtg start codon) [Plasmid F]	(AF013215) ribosomal protein S2 [Bos taurus]	protein kinase C inhibitor-I [Homo	sapiens	Keratin 8 [Homo sapiens]
		_	-							blastx.2		blastx.2		blastx.2		blastx.2	blastx.2	blastx.2		blastx.2
	_									976		226		826		982	586	986		286
										H6EDO51R		HA5AU29R		HACBX26 R		HACMH72	HACMS55 R	HACMX77 HACMX77		HACMZ45
										חלבחחלו	יייי איייייייייייייייייייייייייייייייי	HASAU29		HACBX26	•	HACMH72 HACMH72	HACMS55	HACMX77		HACMZ45

HACND54 HACND54	886	blastx.2	(AC003956)	gb[AAB94632.1]	%19	2	466
			acetolactate synthase		78%	267	404
HACNF21R	991	blastx.2	40-kDa keratin protein	gb AAA36044.1	%98	3	809
HACNF41R	992	blastx.2	cytokeratin 8 (279 AA)	emb CAA31376.1	100%	2	91
			[Homo sapiens]			-	,
HACNI47R	993	blastx.2	ribosomal protein S17	gb AAA60284.1	%56	96	200
HADET44R	966	blastx.2	URF 1 (NADH	emb CAA24026.1	%56	62	481
			dehydrogenase subunit)		92%	481	522
HAHHD12	866	blastx.2	(AF016252)	gb AAB72005.1	%96	F	96
			Spinophilin [Rattus				
HALSG11R	999	blastx.2	(AC004544)	gb AAC12952.1	74%	116	376
			cytochrome C oxidase;		74%	54	134
			match to P14406				
	-		(PID:g117121) [Homo				
		•	sapiens]				
HAOSB87R	1002	blastx.2	delta- aminolevulinate	emb CAA39794.1	73%	-1	549
	-		synthase		-		
			(housekeeping) [Homo			-	
			sapiens		-		_
HAOSE70R	1004	blastx.2	acidic ribosomal	gb AAA36471.1	%88	166	267
			phosphoprotein (P1)		25%	273	359
			[Homo sapiens]				
HAOSF68R	1005	blastx.2	laminin-binding protein	gb AAA36161.1	%06	9/	576
			[Homo sapiens]	••			

452	330	142	233	224	332	405	255	353	303	5	200	379	494	27	343	96	572	429	451		170		13	365	-5-
111	10	2 0	135	135	156	223	223	m	4	†	30	224	336	1	98	4	432	340	7	,			ļ	36.	720
82%	95%	100%	75%	73%	46%	39%	72%	%56	10007	8,001	94%	%96	71%	%88	91%	93%	25%	%08	%86		94%			%96	04½0 T
gb AAA60286.1	gb AAA03081.1	pir JH0302 JH0302						emb CAA26879.1	11 2 4 4 3 5 1 5 5 1 1	gb AAA30133.1	ob[AAA51681.1]				ob[AAD53521,1]AF1	55581			gb AAD27722.1 AF1	32947 1	dbj BAA02991.1			emb CAA39417.1	
ribosomal protein S20	in L7	1_	hornworm (fragments)					lactate dehydrogenase-	A Homo sapiens	Ku protein subunit	LIVING Supreme	o denogrilhomonysteine	aucilosymoniocysicino	nydrolase [Homo	(AF155581)	arotescome subunit	hete 7 Manio reriol		(AF132947) CGI-13	protein [Homo sapiens]	carboxyl	methyltransferase	[Homo sapiens]	HL23 ribosomal	protein [Homo sapiens]
blastx.2	blastx.2	blastx.2					-	blastx.2		blastx.2	talogue 2	Olasta.2			blocks 2	Diasta.4			blastx.2		blastx.2			blastx.2	
1006	1007	1008		-				1009		1011	6101	7101			1012	CIAT			1021		1023			1026	
HAOSG95	HAOSI79R	HAOSJ27R						HAOSJ33R		HAOSK38	K	HAUSLSOK			T. V. T. O. V. T.	HAUSL4/K			HAOTE06R		HAOTF90R			HAOTI07R	
HAOSG95	HAOSI79	HAOSJ27						HAOSJ33		HAOSK38	_	HAOSE36			27, 200	HAUSL47			HAOTE06		HAOTF90			HAOTI07	

80% 361 405	88% 9 356	300		,	97% 57 452	57	57	57	57	57	57	57 1 274 473	57 1 2 274 473 261	57 1 1 274 473 261 146	57 1 1 274 473 261 146	57 1 2 274 473 261 146	57 1 274 473 261 146	2 2 274 473 261 146 106	57 1 1 274 473 261 146 106	274 473 261 146 106 3	57 1 274 473 261 146 106 106	57 2 274 473 261 146 106 106	2 2 274 473 261 146 106 108 3 3 3
					92.1 97%					HU	THO		HTU	2	<u>p</u>	<u>P</u>	<u>P</u>	<u>B</u>	<u> </u>	<u>p</u>	<u>p</u>	<u>P</u>	
sp Q99426 CKAP_H UMAN	í	•		emb CAA30792.1			sp P05787 K2C8_HU	5787 K2C8_HU	5787 K2C8_HU	sp P05787 K2C8_HU MAN dbj BAA07292.1	5787 K2C8_HU I AA07292.1	5787 K2C8_HU I AA07292.1	5787 K2C8_HU I AA07292.1	sp P05787 K2C8_HU MAN dbj BAA07292.1 pir JC1348 JC1348	5787 K2C8_HU I AA07292.1 21348 JC1348	5787 K2C8_HU I AA07292.1 S1348 JC1348 AA11465.1	5787 K2C8_HU I AA07292.1 348 JC1348 AA11465.1	5787 K2C8_HU I AA07292.1 AA11465.1 AC16021.1	5787 K2C8_HU I AA07292.1 AA11465.1 AC16021.1	5787 K2C8_HU I AA07292.1 AA11465.1 AC16021.1 AA91639.1	5787 K2C8_HU I AA07292.1 AA11465.1 AC16021.1 AA91639.1	5787 K2C8_HU I AA07292.1 AA11465.1 AC16021.1 AA91639.1	sp P05787 K2C8_HU MAN dbj BAA07292.1 pir JC1348 JC1348 gb AAC16021.1 gb AAC16021.1 emb CAA79716.1
sp Q99426 UMAN	1			emb CAA3		,	sp P05787 F	sp P05787 k MAN	sp P05787 ^k MAN	sp P05787 8 MAN dbj BAA07	sp P05787 R MAN dbj BAA07:	sp P05787 R MAN dbj BAA07.	sp P05787 R MAN dbj BAA07;	sp P05787 R MAN dbj BAA07: pir JC1348 .	sp P05787 R MAN dbj BAA07.	sp P05787 K2C8 MAN dbj BAA07292.1 pir JC1348 JC134 dbj BAA11465.1	sp P05787 R MAN dbj BAA07; pir JC1348 .	sp P05787 K2C8 MAN dbj BAA07292.1 pir JC1348 JC13 dbj BAA11465.1	sp P05787 R MAN dbj BAA07; pir JC1348 ; dbj BAA114 gb AAC160	sp P05787 K2C8 MAN dbj BAA07292.1 pir JC1348 JC13 dbj BAA11465.1 gb AAA91639.1	sp P05787 R MAN dbj BAA07. dbj BAA114 gb AAC160 gb AAA916	sp P05787 R MAN dbj BAA07; dbj BAA114 gb AAC160 gb AAA916	sp P05787 R MAN dbj BAA07; pir JC1348 - dbj BAA114 gb AAC160 gb AAA916
CYTOSKELETON- ASSOCIATED	(1111)	PROTEIN CKAPI	(TUBULIN FOLDING COFACTOR B).	ME491 antigen	precursor (AA -1 to	237) [Homo saniens]	237) [Homo sapiens] KERATIN, TYPE II	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subumit 1 [Homo	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens]	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens]	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens]	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (KS) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens] CAG-isl 7 [Homo	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens] CAG-isl 7 [Homo sapiens]	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens] CAG-isl 7 [Homo sapiens]	EXT) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens] CAG-isl 7 [Homo sapiens] lumican [Homo sapiens]	237) [Homo sapiens] KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens] CAG-isl 7 [Homo sapiens] lumican [Homo sapiens]	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (KS) (CK 1 cytochrome c oxidase subunit 1 [Homo sapiens] hypothetical 18K protein - goldfish mitochondrion ribosomal protein L39 [Homo sapiens] CAG-isl 7 [Homo sapiens] lumican [Homo sapiens]
	`		ŀ	blastx.2 ME4	precu	blastx.2 KER.		CYT	CYT (CYT	CYT((CYT (K8) blastx.2 cytoc													
1027 61				_		1029 bi	_			1032 51													
HAOTT79R				HAOTU79	~	HAOTW22	~																
HAOTT79				HAOTU79		HAOTW22				HAPNK45					HAPNK45 HAPPR43	HAPNK45 HAPPR43 HAQML40	HAPNK45 HAPNK45 R HAPPR43 HAPPR43R HAQML40 R	HAPNK45 HAPPR43 HAQML40 HAUAK54	HAPNK45 HAPPR43 HAQML40 HAUAK54	HAPPR43 HAQML40 HAUAK54	HAPPR43 HAQML40 HAUAKS4 HAZAC68	HAPNK45 R HAPPR43 HAPPR43R HAUAK54 R HAZAC68 R HAZAC68 R	HAPNK45 HAPNK45 HAPPR43 HAPPR43F HAQML40 R HAUAK54 HAUAK54 R HAZAC68 R HAZAD13 HAZAD13

46% 105 335	105	123	105		70,000		100% 72	100% 72	96% 2 100% 72 97% 1 58% 497	90% 2 100% 72 97% 1 58% 497 81% 579 34% 537	90% 72 100% 72 97% 1 58% 497 81% 579 34% 537	96% 2 100% 72 72 72 87% 497 81% 579 844 637 846 637 946 637 946 637 946 637 946 637 946 637 946 646 646 646 646 646 646 646 646 646	90% 72 100% 72 58% 497 81% 579 34% 537 100% 84	90% 72 72 72 88% 497 81% 579 84 100% 84 81	96% 2 100% 72 2 58% 497 81% 579 34% 537 100% 84
46	40	38	200		emb CAA62211.1 96			HT 1							
					emb CAA62		sp P05787 K MAN	sp P05787 K MAN	sp P05787 K MAN gb AAC526	sp P05787 K MAN gb AAC526					
fibronectin precursor	Target and the control of			•	unnamed protein product [Homo	sapiens	KERATIN, TYPE II CYTOSKELETAL 8	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N-	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N- methyltransferase [Rattus norvegicus]	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N- methyltransferase [Rattus norvegicus] (AF064205) dynactin 1	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N- methyltransferase [Rattus norvegicus] (AF064205) dynactin 1 p135 isoform [Homo saniens]	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N- methyltransferase [Rattus norvegicus] (AF064205) dynactin 1 p135 isoform [Homo sapiens] (AL031427) dJ167A19.3 (novel protein) [Homo	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N- methyltransferase [Rattus norvegicus] (AF064205) dynactin 1 p135 isoform [Homo sapiens] (AL031427) dJ167A19.3 (novel protein) [Homo sapiens]	KERATIN, TYPE II CYTOSKELETAL 8 (CYTOKERATIN 8) (K8) (CK 1 protein arginine N- methyltransferase [Rattus norvegicus] (AF064205) dynactin 1 p135 isoform [Homo sapiens] (AL031427) dJ167A19.3 (novel protein) [Homo sapiens]
	-				blastx.2	ļ	blastx.2	blastx.2	blastx.2 blastx.2	blastx.2 blastx.2	blastx.2 blastx.2 blastx.2 blastx.2	blastx.2 blastx.2 blastx.2 blastx.2	blastx.2 blastx.2 blastx.2 blastx.2 blastx.2	blastx.2 blastx.2 blastx.2 blastx.2 blastx.2	blastx.2 blastx.2 blastx.2 blastx.2 blastx.2
					HAZAG23 1046 R	+	HAZA189K 1051				2R 2R		2R 2R 80	2R 2R 9R	9R 80 2R
HAZAE44 HAZAE44K					HAZAG23 HAZ	十	HAZALIOY LITA					0	0	0	0

				[Oryctolagus cuniculus]				
HBJHY72	HBJHY72R	1063	blastx.2	(AL117434)	emb CAB55922.1	%86		297
:	P00B			hypothetical protein				
	_			[Homo sapiens]				
HBXCG52		1065	blastx.2	(AK001079) unnamed	dbj BAA91496.1	%68	371	427
	R			protein product [Homo sapiens]		46%	457	909
HCACS53	HCACS53R	1067	blastx.2	histone H3 [Spisula	gb AAA29965.1	%06	313	507
	POOA			solidissima]		100%	116	244
202 12022						%98	206	574
HCHAJ85	HCHAJ85R	1068	blastx.2	hemolysin	gb AAA58585.1	61%	206	57
				[Acanthamoeba				
				polyphaga]				
HCHIMIM7	HCHIMIM71	1069	blastx.2	(AC003040) unknown	gb[AAC23757.1]	28%	2	244
4	≭		. '	protein [Arabidopsis		44%	249	329
, 01. 6				Liamana				
HCLBH21	HCLBH21R	1071	blastx.2	cytochrome c oxidase	dbj BAA85270.1	%16	3	416
				subunit 1 [Pan		%88 88	415	009
HCOMA72	HCOMA72 HCOMA72	1073	blastx.2	NuMA protein - human	pir S33413 S33413	100%	115	273
TOOM OUT	TICONO	1074	1.1.					
toomport		10/4	olastx.2	prolylcarboxypeptidase	gb AAA99891.1	%86	77	718
2001	N.		,	Homo sapiens		25%	702	782
HCOMD38 HCOMD3	HCOMD38	1075	blastx.2	phospholipid	emb CAA50793.1	100%	39	275
	4			nydroperoxide			_	
			_	glutathione peroxidase				-
				Homo sapiens				-
HCOMD61	HCOMD61	1076	blastx.2	(AF047470) malate	gb[AAC03787.1]	81%	-	216

287		655	522 98		046	401	1		240	301	268						470		176			282		
240		4.	100 54		35	215	CtC		304	68	518						273		738			4		
75%		%08	71:%		- %9L	7050	2		. 59%	%99	64%				,		-% 6		100%			%86		
		gb AAB03345.1	gb AAA18502.1		emb CAA55026.1	11 4 A DOODON 11	go AAD0000/.1		emb CAA86061.1								emb CAA71575.1	-	dbj BAA03853.1			gb AAC05826.1		
dehydrogenase	precursor [Homo sapiens]	hypothetical protein 384D8_6 [Homo	elongation factor 1 alpha [Oryctolagus	cuniculus	ribosomal protein L15	Kattus norvegicus	ZZ:beta-Gal' 1gG- binding fision protein	funidentified cloning 1	similar to 40S	ribosomal protein;	cDNA EST	CEMSA13F comes	from 1 1 gene; cDNA	EST EMBL:M79582	comes from this gene;	cDN	fused-ccdB	[Escherichia coli]	The polymorphysm	(RFLP) of this gene is	associated with 1 1	(AF042857) lung	cancer antigen NY-LU-	12 variant A [Homo
		blastx.2	blastx.2		blastx.2		blastx.2		blastx.2		•						blastx.2		blastx.2			blastx.2		
		1078	1079		1080		1083		1084								1085		1087			1088		
R		HCOMF52 R	HCOMG28	4	HCOMG40	R	HCOMI30R		HCOMT37R								HCOML11	24	HCOMM55	24		HCOMO58	R	
		HCOMF52	HCOMG28		HCOMG40 HCOMG40		HCOMI30		HCOMI37) CITATOOTT							HCOME 11		HCOMMS	5		HCOMO58		

		Ĺ		sapiens				
HCOMW5	HCOMW52 RP00B	1089	blastx.2	(AB032025) ubiquitin [Canis familiaris]	dbj BAA83996.1	%66	09	443
HCOMX77		1090	blastx.2	HETEROGENEOUS	sp P09651 ROA1_H	%59	133	315
	ᅺ			NUCLEAR	UMAN	%96	3	82
 -			·	RIBONUCLEOPROTE		100%	81	122
				N A11 (HNRNP		. 57%	٠,	82
				CORE PROTEIN A1).		53%	∞	92
						48%	5	82
				÷		34%	142	291
						27%	11	82
						62%	81	128
						44%	٧,	82
						45%	=======================================	79
		•	<u>.</u>	-		75%	81	116
						20%	84	119
HCONCIS	HCONC18	1091	blastx.2	cathepsin D [Homo	gb AAA51922.1	%18	3	188
	K-00B			sapiens]		91%	190	333
						83%	335	424
						71%	426	467
HCONKSO HCONK	HCONKS6	1093	blastx.2	ribosomal protein L18	gb AAA16329.1	100%	104	616
	Y.			[Homo sapiens]		27%	267	999
	HCONL49 R	1095	blastx.2	ribosomal protein S13 [Homo sapiens]	dbj BAA13528.1	%86	m	221
HCONO17	HCONO17	1099	blastx.2	(AC004240) match to	gb AAC04502.1	81%	4	69
	4			Z43555 (NID:g5/2/88) [Homo sapiens]		71%	65	148
HCON025	HCONO25	1100	blastx.2	5,10-	gb AAC41945.1	77%	251	610
	~			methenyltetrahydrofola	-	87%	147	245
				te synthetase [Homo	-	100%	m	89

H+-transporting ATP synthase (EC 3.6.1.34) 58K chain - human
1.2.1.12) [Homo
- 1
fruman tenacycure
transporter-like proteint mRNA [Homo sapiens]
ornithine decarboxylase
sapiens
calpactin I light chain

		439	470	652	_	\$15	406	748	552	310	365		-	426						367	573	
		95	171	473	_	30	233	869	49	218	330			22			•	,	<u> </u>	11	367	
	•	100%	%66	77%		25%	62%	52%	%82	87%	83%	•		91%	<u></u> _					%86	62%	
		gb AAA35646.1	gb[AAC69149.1]	-	·	gb AAD33912.1 AF1	43815_1		gb AAA59461.1	gb AAD25980.1 AF0	95770_1			sp P32119 TDX1_HU MAN						pir A56846 A56846		
nhosnhatasa (Homo	sapiens]	neutral protease alpha subunit [Homo sapiens]	(AF019661) zeta	proteasome chain;	PSMA5 [Mus	(AF143815) ribosomal	protein [Bos taurus]		keratin 18 [Homo sapiens]	(AF095770) PTH-	responsive	osteosarcoma D1	protein [Homo sapiens]	THIOREDOXIN PEROXIDASE 1	(THIOREDOXIN- DEPENDENT	PEROXIDE 1	(NATURAL KILLER	CELL ENHANCING	FACTOR B) (NREF-B).	ribosomal protein L19 -	rat	٠.
		blastx.2	blastx.2			blastx.2			blastx.2	blastx.2				blastx.2						blastx.2		
		1120	1121			1122			1123	1126				1127		•				1129		
R		HCOOT43 R	HCOOT68	X		HCOOUS6	8		HCOOW72 R	HCOOX48	~			HCOOY43				,	_	HCOPB03R		
		HCOOT43	HCOOT68			HC00US6		- 1	HC00W7	HC00X48				HC00Y43						HCOPB03		

							0.10	
68	541	400	310	337	507	593	579 625	
3	104	2	62	122	4	3	572	
%96	91%	%66	77%	%59	74%	%26	%£8 %96	
emb CAA34890.1	gb AAA31492.1	gb AAB46780.1	emb CAA67127.1	gb AAC52076.1	dbj BAA03401.1	emb CAA71256.1	emb CAA20237.1	
homologue of yeast IPP isomerase [Homo sapiens]	ubiquitin conjugating- protein [Oryctolagus cuniculus]	serine protease homolog=NES1 [human, mammary epithelial cells, 76N, Peptide, 276 aa] [Homo sapiens]	fused; toxic gene [synthetic construct]	(AF026246) HERV-E envelope glycoprotein Homo sapiens]	'human homologue of rat ribosomal protein L9' [Homo sapiens]	MEMD protein [Homo sapiens]	(AL031228) dJ1033B10.9 (Short- chain alcohol	dehydrogenase family member (HKE6, RING2)) [Homo
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	·
1130	1131	1132	1134	1136	1138	1139	1140	
HCOPC45R	HCOPD67R	HCOPE27R	HCOPI09R A	HCOPO34R	HCOPO88R	HCOPV41R	HCOPZ15R	
HCOPC45	НСОРД67	HCOPE27	HCOPI09	HCOPO34	HCOPO88	HCOPV41	HCOPZ15	

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				1,000				
	_			sapieris				
HC0QA38	HCOQA38 R	1141	blastx.2	hBD-1 [Homo sapiens]	emb CAA63405.1	100%	89	271
HCOQB12	HC0QB12	1143	blastx.2	(AC005600) tuberin	gb AAC34210.1	100%	4	309
,	R			[Homo sapiens]		84%	308	346
HC0QD29	нсоор29	1147	blastx.2	S3 ribosomal protein	gb AAB19349.1	93%	16	588
,	8			[human, colon, Peptide,		25%	552	707
				243 aa] [Homo sapiens]	٠	%09	632	730
нсоорзя	HCOQD38	1148	blastx.2	HMG1 protein (AA 1 - 215) [Bos taurus]	emb CAA31284.1	%56	112	989
HC00D49	HCOQD49	1149	blastx.2	H+-transporting ATP	pir A32019 PWBOG	%68	35	532
,				synthase (EC 3.6.1.34)		%59	513	692
				gamma chain precursor			,	
				- povine				
HCOQG37	HC0QG37	1151	blastx.2	1-8D [Homo sapiens]	emb CAA40625.1	866	51	386
	R					100%	386	409
нсоон46 нсоон	нсоон46	1153	blastx.2	mucin MUC5B [Homo	gb AAC51343.1	100%	95	343
	ጸ			sapiens]		64%	324	365
HCOQJÖ6	HCOQJ06R	1156	blastx.2	37kD Laminin receptor	emb CAA64147.1	85%	56	361
				precursor /p40		28%	352	474
•				ribosomal associated				
				protein [Gallus gallus]				
HC0Q107	HCOQ107R	1157	blastx.2	laminin-binding protein	gb AAA36161.1	%88	.9	959
				[Homo sapiens]				
HCOQJ79	HCOQJ79R	1159	blastx.2	elongation factor 1	gb[AAA18502.1]	83%		471
				alpha [Oryctolagus				
	•			cuniculus]				
HCOQK86 HCOQK	HCOQK86	1160	blastx.2	(AF038129)	gb AAB92373.1	%46	4	456
,	~			polyubiquitin [Ovis		94%	4	456

				aries]	٠	93%	28	456
						%26	4	255
					· ·	26%	440	487
						30%	440	556
		:				30%	440	556
HCOQL87	HCOQL87	1161	blastx.2	glutathione S-	gb AAA56823.1	%95	24	521
	~			transferase-pi [Homo		41%	328	630
HCOQM87	нсоомял нсоомял	1162	blastx.2	DRPLA protein [Mus	dbj BAA13450.1	52%	m	278
	R			musculus]		47%	17	307
HCOQO79	HC00079	1163	blastx.2	TIMP [Homo sapiens]	emb CAA00898.1	%66	2	574
	R					61%	561	809
HCOQP32	HCOQP32R	1164	blastx.2	elongation factor 1-	gb AAA52367.1	%56	15	149
				alpha [Homo sapiens]		64%	151	381
HCOQS11	HCOQS11R	1165	blastx.2	acidic ribosomal	gb AAA36472.1	78%	102	398
				phosphoprotein (P2)				_
				TYOUTO SUPPOSE				
НСОФО92	HCOQU92 R	1167	blastx.2	protein translocation	gb AAA19639.1	100%	91	270
				[Canis familiaris]		2001	1	100
HCOQV27 HCOQV27	HCOQV27	1168	blastx.2	B4-2 protein [Homo	gb AAA85576.1	%96	3	239
	R			sapiens]				
HCOQX38	нсоох38	1169	blastx.2	catechol O-	emb CAA81263.1	%96	32	421
	પ્ર			methyltransferase		91%	421	489
				[Homo sapiens]		%08	495	539
						100%	П	30
НСООУЗЗ	HCOQY33 R	1170	blastx.2	(AB000910) ribosomal protein [Sus scrofa]	dbj BAA19210.1	%96	49	321
HCOQZ86	HCOOZ86	1173	blastx.2	MHC class II HLA-	gb AAA59782.1	92%	106	357

	т								_							,					
470	292	482	317	305	305	305	317	317	124	419	245	276	369	416	427	463	408	136	409	293	470
351	89	75	120	126	123	129	120	132	2	3	3	247	1	297	332	38	115	38	326	273	255
77%	%86	94%	100%	.007 26%	46%	20%	44%	43%	39%	%86	91%	%06	87%	26%	47%	%16	61%	%06	23%	100%	%26
	dbj BAA28169.1	gb AAA85657.1	dbj BAA06031.1				•			emb CAA40655.1	dbj BAA77672.1		gb AAC48728.1			gb AAA03341.1	gb AAF17196.1 AF1	12208_1			gb AAA91461.1
DR-beta-1 [Homo sapiens]	(AB012122) TIP49 [Homo sapiens]	ribosomal protein L28 [Homo sapiens]	hnRNP B1 protein	Litolilo sapielis						pm5 protein [Homo sapiens]	NADH dehydrogenase	subunit 3 [Homo sapiens]	IGF binding protein-2	[Sus scrofa]		ribosomal protein L23a [Homo sapiens]	(AF112208) 13kDa	differentiation-	associated protein	[Homo sapiens]	UbcHSC [Homo sapiens]
	blastx.2	blastx.2	blastx.2			,				blastx.2	blastx.2		blastx.2			blastx.2	blastx.2				blastx.2
	1175	1177	1178							1179	1180		1181			1182	1185		٠		1186
R	HCORB20 R		HCORI18R							HCORI25R	HCQCR82	R	HCRME42	~		HDABR53 R	HDTAQ74	&			HDTBP08R
	HCORB20	HCORB66	HCORI18								HCQCR82		HCRME42 HCRME42			HDABR53	HDTAQ74				HDTBP08

275	441	288	475	549		170		376		. <u> </u>		128			179	321		130	284		406		517			325	
27	259	49	272	475	1	m		62				m			m	178		7	186		152		7			2	
%68	%88	93%	95%	%08		100%		37%		<u></u> -		100%			%68	%68			% 69		%96		93%			%06	
dbj BAA85270.1		gb AAD01439.1				emb CAA59735.1		gb[AAD15346.1]				pir S12206 S12206			emb CAA24033.1			dbj BAA07291.1			gb[AAA97434.1]		gb AAA20843.1			emb CAA24033.1	
cytochrome c oxidase	subunit 1 [Pan troglodytes]	(AF010472) alpha-	amidating	monooxygenase [Homo	sapiens	alligator Wilm's tumour	protein [Alligator	(AC004044) predicted	protein of unknown	function [Arabidopsis	thaliana]	hypothetical protein 2	(rRNA external	transcribed spacer) - 1	URF 3 (NADH	dehydrogenase subunit)	[Homo sapiens]	NADH dehydrogenase	subunit 2 [Homo	sapiens	cytokine SDF-1-beta	[Homo sapiens]	cytochrome oxidase	subunit II [Homo	sapiens]	URF 3 (NADH	dehydrogenase subunit)
blastx.2		blastx.2				blastx.2		blastx.2				blastx.2			blastx.2			blastx.2			blastx.2		blastx.2			blastx.2	٠
1188		1190		•		1191		1192				1193			1194			1196			1198		1199			1200	
HDTDB88	24	HE8QX44R	,			HE9QU94R		HEAHF02R				HEEAY40R			HEGAF68R			HFABK01R			HFIBG63R		HFIJB15R			HFIXK57R	
HDTDB88		HE8QX44	,			HE9QU94		HEAHF02				HEEAY40			HEGAF68			HFABK01			HFIBG63		HFIJB15			HFIXK57	

				[Homo sapiens]				
HFIZQ64	HFIZQ64R	1201	blastx.2	(AK001601) unnamed protein product [Homo	dbj BAA91782.1	100%	48	137
				sapiens]				
HFKKK36	HFKKK36 R	1202	blastx.2	neutral calponin [Homo sapiens]	dbj BAA12090.1	%96		98
HFPEC93	HFPEC93R	1203	blastx.2	URF 2 (NADH	emb CAA24027.1	87%	99	527
				dehydrogenase subunit)		%06	П	99
HFPIX37	HFPIX37R	1204	blastx.2	(AL137696)	emb CAB70878.1	51%	4	240
				hypothetical protein		41%		258
HFTDK36	HFTDK36R	1205	blastx.2	folate-binding protein	ph/AA35822 11	100%	06	000
				precursor [Homo		38%	341	514
		,		sapiens]			_	-
HFVIB28	HFVIB28R.	1206	blastx.2	fused-ccdB	emb CAA71575.1	%02	56	175
				[Escherichia coli]				
HFXGR60	HFXGR60R	1207	blastx.2	(AF034746) LNXp70	gb AAC40076.1	%95	347	57.
200000000000000000000000000000000000000	2000111111	900		Sunacutt sulvi				
HHAUD07		1208	blastx.2	GIP-binding	pir JH0813 JH0813	15%	337	459
	ፈ			regulatory protein Gs alpha chain isoform - 1				-
HHBF021	HHBF021R	1209	blastx.2	URF 3 (NADH	emb CAA24033.1	83%	3	236
•				dehydrogenase subunit)		85%	248	328
				[Homo sapiens]		64%	205	246
HHEVG50	HHEVG50	1210	blastx.2	CAG-isl 7 [Homo	gb AAC16021.1	93%	13	474
\neg	R			sapiens]				
HHFGQ65	ннгооб	1211	blastx.2	(AJ249731) putative	emb CAB56506.1	81%	107	319
	RA			G8.1 protein Homo		77%	25	129

888) NADH gb AAC25447.1 78% 3 genase subunit 0. sapiens 3 o sapiens emb CAB71165.1 74% 2 III [Talpa gb AAA19775.1 89% 79 al ab 324 bane b [Homo gb AAA19775.1 84% 61 B2072 [Homo gb AAA31002.1 100% 3 Homo sapiens] gb AAA59461.1 100% 3 [Homo sapiens] gb AAA59461.1 100% 2 [Homo sapiens] gb AAA63538.1 100% 3 chochty [Homo land) schlas submit, 20 3 chelta submit, 20S [Homo land) 100% 1 chelta submit, 20S [Homo land) 1 100% 1 chelta submit, 20S [Homo land) 1 1				-	Consisted			-	
1214 blastx.2 cytochrome oxidase emb CAB71165.1 74% 2 2 [Homo sapiens] emb CAB71165.1 89% 79 2 [Homo sapiens] emb CAB71165.1 89% 79 2 [Homo sapiens] emb CAB7116.1 84% 61 3 [Homo sapiens] emb CAB7116.1 100% 3 3 [Homo sapiens] emb CAB7116.1 100% 3 4 [Homo sapiens] emb CAB7116.1 100% 3 5 [Homo sapiens] emb CAB7116.1 100% 19 5 [Homo sapiens] emb CAB7116.1 100% 19 6 [Homo sapiens] emb CAB7116.1 100% 1 6 [Homo sapiens] emb CAB7116.1 100% 1 7 [Homo sapiens] emb CAB7116.1 100% 1 7 [Homo sapiens] emb CABA17675.1 98% 1 7 [Homo sapiens] emb CABA17675.1 98% 1 7 [Homo sapiens] emb CABA17675.1 98% 1 8 [Homo sapiens] emb CABA17675.1 98% 1 9 [Homo sapiens] emb CABA17675.1 98% 1	H	SEC15B	1213	blacty 2	(AF014888) NADH	gblAAC25447.1	78%	3	743
1214 blastx.2 2 Iroliu sapiens 1215 blastx.2 cytochrome oxidase emb CAB71165.1 74% 2 170 2 subunit II [Talpa europaeal 1215 blastx.2 cytochrome b [Homo gb AAA19775.1 89% 79 79 79 79 79 79 79	Ē	ACIO TO	6171	oldasio.	dehydrogenase subunit				
Subunit III [Talpa europaea] 1215 blastx.2 cytochrome b [Homo gb AAA19775.1 89% 79 79 1216 blastx.2 (AB015335) HRIHFB2072 [Homo sapiens] 1218 blastx.2 ATP synthase beta subunit precursor [Sus scrofa] 1220 blastx.2 complex delta subunit, partial CDS [Homo gb AAA51809.1 100% 19 1221 blastx.2 ubiquitin-conjugating emb CAA63538.1 100% 19 1222 blastx.2 complex delta subunit, partial CDS [Homo gb AAA51809.1 100% 1 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1	臣	ISGP15R	1214	blastx.2	cytochrome oxidase	emb CAB71165.1	74%	2	529
1215 blastx.2 cytochrome b [Homo gb AAA19775.1 89% 79 79 79 79 79 79 79		7			subunit III [Talpa	-	. 62%	517	717
1215 blastx.2 cytochrome b [Homo gb AAA19775.1 89% 79 79 79 79 79 79 79					europaea]				
1216 blastx.2 (AB015335) dbj BAA88116.1 84% 61 1218 blastx.2 HRIHFB2072 [Homo sapiens] 1220 blastx.2 sapiens] 1221 blastx.2 sapiens 1222 blastx.2 sapiens 1222 blastx.2 sapiens 1222 blastx.2 sapiens 1224 blastx.2 sapiens 1224 blastx.2 sapiens 1224 blastx.2 sapiens 1225 sapiens 1	田と	ISGQ17	1215	blastx.2	cytochrome b [Homo sapiens]	gb AAA19775.1	86%	79	435
HRIHFB2072 [Homo sapiens] 100% 324 1218 blastx.2 Na+, K+-ATPase beta subunit precursor [Sus scrofa] 1220 blastx.2 sapiens] 1221 blastx.2 antigen [Homo sapiens] 1222 blastx.2 antigen [Homo sapiens] 1223 blastx.2 antigen [Homo sapiens] 1224 blastx.2 ATP synthase beta subunit precursor 1225 blastx.2 ATP synthase beta 1225 blastx.2 ATP synthase beta 1225 blastx.2 antipopontin [Homo sapiens] abject 1225 antipopontin [Homo sapiens] abject 1225 antipopontin [Homo sapiens] abject 1225 antipopontin [Homo sapiens] abject antipopontin [Homo sapiens] antipopontin [旧	CBAD05	1216	blastx.2	(AB015335)	dbj BAA88116.1	84%	61	351
1218 blastx.2 Na+, K+-ATPase beta-subunit precursor [Sus scrofa] 100% 1 1020 blastx.2 keratin 18 [Homo sapiens] 221 blastx.2 complex delta subunit, partial CDS [Homo 1224 blastx.2 ATP synthase beta subunit precursor 224 blastx.2 ATP synthase beta subunit 225 blastx.2 ATP synthase beta 226 AAA51809.1 100% 1 100%	K				HRIHFB2072 [Homo		100%	324	356
1218 blastx.2 Na+, K+-ATPase beta-subunit precursor [Sus subunit precursor [Sus scrofa] 100% 1 1020 blastx.2 keratin 18 [Homo apiens] 221 blastx.2 antigen [Homo sapiens] 222 blastx.2 blastx.2 complex delta subunit, partial CDS [Homo 1 1224 blastx.2 ATP synthase beta 225 blastx.2 ATP synthase beta 226 AAA51809.1 100% 1 1224 blastx.2 ATP synthase beta 225 blastx.2 Camplex delta subunit precursor 225 blastx.2 225 blastx.2 225 blastx.2 225 blastx.2 225 blastx.2 225 blastx.2 225 blastx.3 225					sapiens]				
Subunit precursor [Sus scrofa] Scrofa] Scrofa Scrofa Scrofa Scrofa Sapiens Sapie	呂	KZAE07R	1218	blastx.2	Na+, K+-ATPase beta-	gb AAA31002.1	100%		138
1220 blastx.2 keratin 18 [Homo gb AAA59461.1 100% 3 1221 blastx.2 antigen [Homo sapiens] gb AAA02999.1 100% 2 1222 blastx.2 ubiquitin-conjugating emb CAA63538.1 100% 19 1223 blastx.2 (AC005545) AP-3 gb AAC34214.1 98% 3 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 1225 blastx.2 ATP synthase beta subunit precursor Homo sapiens] Homo sapiens] 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1					subunit precursor [Sus				
1221 blastx.2 antigen [Homo sapiens] gb AAA02999.1 100% 2 1222 blastx.2 ubiquitin-conjugating emb CAA63538.1 100% 19 1223 blastx.2 (AC005545) AP-3 gb AAC34214.1 98% 3 1224 blastx.2 ATP synthase beta subunit, partial CDS [Homo 1 subunit precursor Homo sapiens] 1225 blastx.2 homo sapiens] 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1	田	KZAI14R	1220	blastx.2	keratin 18 [Homo	gb AAA59461.1	100%	æ	368
1221 blastx.2 antigen [Homo sapiens] gb AAA02999.1 100% 2 1222 blastx.2 ubiquitin-conjugating emb CAA63538.1 100% 19 1223 blastx.2 (AC005545) AP-3 gb AAC34214.1 98% 3 1224 blastx.2 ATP synthase beta subunit precursor Homo sapiens] 1225 blastx.2 homo sapiens] Homo sapiens		_			sapiens				i
1222 blastx.2 ubiquitin-conjugating emb CAA63538.1 100% 19 1022 enzyme UbcH7 [Homo 1223 blastx.2 (AC005545) AP-3 complex delta subunit, partial CDS [Homo 1 partial CDS [Homo 1 subunit precursor Homo sapiens] Homo sapiens] Electrical Complex delta subunit precursor Homo sapiens] Homo sapiens Hom	H	KZAI68R	1221	blastx.2	antigen [Homo sapiens]	gb AAA02999.1	100%	2	79
complex OboH7 [Homo sapiens] sapiens sapiens sapiens sapiens complex delta subunit, partial CDS [Homo 1 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 Homo sapiens Homo sapiens Sb AAA17675.1 98% 1 Homo sapiens 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1 Homo sapiens 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1 Homo sapiens 1225 blastx.2 nephropontin [Homo gb AAA17675.1 1225 blastx.2 nephropontin [Homo gb AAA17675.1 1 1 1 1 1 1 1 1 1	H	KZAQ39	1222	blastx.2	ubiquitin-conjugating	emb CAA63538.1	100%	19	480
1223 blastx.2 (AC005545) AP-3 gb AAC34214.1 98% 3 complex delta subunit, partial CDS [Homo 1 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 lhomo sapiens] Homo sapiens] gb AAA17675.1 98% 1 lhomo sapiens	24				enzyme UbcH7 [Homo sapiens]				
complex delta subunit, partial CDS [Homo 1 partial CDS [Homo 1 100% 1 100% 1 1224 blastx.2 ATP synthase beta gb AAA51809.1 100% 1 Example Interpretation	田	KZAR58	1223	blastx.2	(AC005545) AP-3	gb AAC34214.1	%86	e.	566
\$59R 1224 blastx.2 ATP synthase beta subunit precursor gb AAA51809.1 100% 1 Homo sapiens [Homo sapiens] [Homo sapiens] 1025 blastx.2 nephropontin [Homo Bb AAA17675.1 98% 1	R .				complex delta subunit, partial CDS [Homo 1				
Subunit precursor Homo sapiens S64R 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1	臣	KZAS59R	1224	blastx.2	ATP synthase beta	gb AAA51809.1	100%		462
Homo sapiens S64R 1225 blastx.2 nephropontin [Homo gb AAA17675.1 98% 1					subunit precursor				
S64R 1225 blastx.2 nephropontin Homo gb AAA1/6/5.1 96% 1			_		Homo sapiens		/000	-	270
	二	KZAS64R	_	blastx.2	nephropontin Homo	gb AAA1/6/5.1	78%	1	0/7

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HK7 A S84	TIV7 A COAD	1006	4-	Saprottal	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		ľ	
	NI-OCUTATIVE I	777	0143tA:4	subunit (EC 3.4.22.17)	go AAA30422.1 	%00I	7	103
				Bos taurus]		÷		
HKZAV69	HKZAV69	1227	blastx.2	ribosomal protein S17	gb AAA60284.1	100%	8	412
	R			[Homo sapiens]	-			
HKZAV72	HKZAV72	1228	blastx.2	fibronectin precursor	emb CAA26536.1	100%	2	163
	R			[Homo sapiens]			•	
HKZBB22	HKZBB22R	1230	blastx.2	(AB015610) ribosomal	dbj BAA36501.1	100%	8	520
				protein S4X				
				[Chlorocebus aethiops]				
HKZBS89	HKZBS89R	1231	blastx.2	ferritin heavy subunit	gb A'AA35830.1	100%	-	381
				[Homo sapiens]	-		ı	
HLDQQ80	нгроово	1232	blastx.2	ATPase subunit 6	emb CAA76999.1	74%	2	142
				[Papio hamadryas]	•	54%	153	257
HLDXE19	HLDXE19R	1233	blastx.2	neuron-restrictive	gb AAC50115.1	%08	17	187
				silencer factor [Homo				
				sapiens]				
HLICD55	HLICDSSR	1234	blastx.2	glutamine:fructose-6-	gb AAA58502.1	%96	56	148
				phosphate				
				amidotransferase				
				[Homo sapiens]				_
HLJB137	HLJB137R	1236	blastx.2	H(+)-transporting ATP	emb CAA45865.1	100%	165	335
				synthase [Bos taurus]		%59	329	610
			,			%08	83	187
						%09	481	555
十		3				%88	450	476
HLIHA4/	HL1HA47K	1237	blastx.2	100 kDa protein	emb CAA45756.1	82%	187	393
				Rattus norvegicus	-	88%	380	406

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subunit)	subunit)	subunit) Idase bates bates ogenase	subunit) ldase bates ogenase l	nit)	nit)	nit)	nit)	nnit)	nit)	nuit)
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	242 70	1 245 315 244	224	52	114	3 509		86	101
	74% 55%	100% 41% 46% 35%	%16	%56	91%	83% 81%	93%	%86	100%
	gb AAC52058.1	gb AAC39523.1	emb CAB43181.1	gb AAD05421.1	gb AAA36471.1	gb AAA30032.1	emb CAB42187.1	emb CAA61582.1	emb CAA49189.1
heavy chain isoform SM1 [human, umbilical 1	growth-arrest-specific protein 2 [Homo sapiens]	OS9 [Homo sapiens]	(AL031670) dJ681N20.2 (similar to FTLL1(ferritin, light 1	(AF044957) NADH:ubiquinone oxidoreductase B15 subunit [Homo sapiens]	acidic ribosomal phosphoprotein (P1) [Homo sapiens]	actin 2 protein [Strongylocentrotus purpuratus]	unnamed protein product [unidentified]	ribosomal protein L21 [Homo sapiens]	ribosomal protein L26 [Homo sapiens]
,	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	1250	1251	1252	1253	1254	1257	1259	1260	1261
ጸ	HNBVL57 R	HNHBC18 R	HNJFE85R	HNKCO29 R	HNOAA22 R	HNOAB88 R	HNOAC15 R	HNOAE50 R	HNOAE65 R
	HNBVL57	HNHBC18	HNJFE85	HNKCO29	HNOAA22	HNOAB88	HNOAC15	HNOAE50	HNOAE65 HNOAE65

233	239	0	319	239	313	239	236	236	236	239	239	236	236	239	236	239	236	239	239	239	319	239	236	239	200	236	239	239
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100%	64%	%7/	75%	37%	%29	36%	44%	36%	37%	44%	39%	42%	38%	38%	45%	48%	35%	41%	35%	36%	62%	45%	47%	36%	38%	45%	35%	38%
emb CAA61582.1	pir A91193 CGBO1S																											
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blastx.2	blastx.2				•																							
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239	236	239	236	236	239	239	322	239	236	239	239	239	319	239	322	. 236	328	328	322	239	319	236	322	316	239	316	316	328	328
93	9	m	69	9	15	84	218	9	21	15	m	n	218	54	227	15	233	233	227	111	218	21	233	233	114	233	233	227	227
44%	41%	37%	41%	35%	34%	44%	48%	35%	36%	33%	36%	32%	48%	36%	\$2%	35%	20%	24%	51%	46%	47%	33%	23%	21%	45%	21%	53%	20%	51%
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233	227	233	227	233	233	227	242	212	227	233	1	230	-	209	260	227	-	_	227	-		-	233	221	-	. 227	-	1	233
20%	51%	20%	41%	20%	46%	46%	44%	41%	46%	46%	45%	44%	36%	42%	63%	45%	44%	41%	43%	45%	45%	44%	46%	36%	45%	45%	41%	38%	46%
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322	316	108	114	108	. 108	316	322	108	108	328	316	316	322	316	105	108	316	93	316	93	316	316	316	93	83	108	316	316	105
233	221	177				212	221			233	227	227	233	221	,1		227	-	227		221	227	227		-		218	260	1
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		elongation factor 1 alpha [Oryctolagus	(AB000911) ribosomal protein [Sus scrofa]	(AD001528)	spermidine	aminopropyltransferase [Homo sapiens]	MAP kinase kinase	Q1Z 7F5 [Homo sapiens]	epoxide hydrolase	TAXREB107 [Homo	(AF161378) HSPC260	[Homo sapiens]	scar protein [Homo	sapiens]	proteoglycan core protein [Homo sapiens]	glutathione transferase M1 [Homo sapiens]	(AB016193) transcription factor
		blastx.2	blastx.2	blastx.2			blastx.2	blastx.2	blastx.2	blastx:2	blastx.2		blastx.2		blastx.2	blastx.2	blastx.2
		1266	1267	1270			1273	1274	1275	1276	1278		1279		1280	1281	1283
		HNOAH67 R	HNOAJ67R	HNOAL51	&		HNOA071 R	HNOAP21 R	HNOAQ24 R	HNOAQ47	HNOAR85	ጸ	HNOAS07	R	HNOAS36 R	HNOAS92 R	HNOAT76 HNOAT76 I.
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general transcription factor [Homo sapiens]	(AF110731)	antioxidant enzyme B166 [Homo sapiens]	midkine [Homo	(AF077054) unr protein [Homo sapiens]	ك-	adenosylhomocysteine hydrolase [Homo	sapiens	ubiquitin [synthetic construct]	(AF035718)	mesoderm-specific	basic-helix-loop-helix	protein; Pod-1 [Homo	sapiens]	ribosomal protein L23a [Homo sapiens]	plasma gelsolin [Homo	sapiens	von Willebrand factor	prepropeptide [Homo	sapiens]			
blastx.2	blastx.2		blastx.2	blastx.2	blastx.2			blastx.2	blastx.2					blastx.2	blastx.2		blastx.2					
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				gb AAA37026.1	1 0000	emb[CAA62188.1]																							
				alpha-tubulin III	[Cricetulus griseus]	Protein sequence and	annotation available	allitoration available	SOON VIA DWISS-1100;	[Homo sapiens]																			
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											-				envelope protein	[Homo sapiens]	(AJ224875)	glucosyltransferase	[Homo sapiens]	signal recognition	particle subunit 14	[Homo sapiens]	hnRNPcore protein A1	[Homo sapiens]	(AF009368) Luman	[Homo sapiens]	ribosomal protein L6	[Homo sapiens]	ribosomal protein L29
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	gb AAC31959.1	gb AAA88855.1	gb AAA52678.1	emb CAA57650.1	gb AAA42074.1	dbj BAA01980.1	emb CAA20229.1		gb AAF07341.1 AF1 96481_1	dbj BAA03401.1	emb CAA00898.1	gb AAF03750.1 AF1
[Homo saniens]	(AF081484) alpha- tubulin isoform 1	MHC class I A [Pan troglodytes]	high mobility group protein 17 [Homo sapiens]	hevin [Homo sapiens]	ribosomal protein L5 [Rattus norvegicus]	ORF [Pan troglodytes]	(AL031228) dJ1033B10.2 (WD40 protein BING4 (similar	to S. 1 sapiens]	(AF196481) RING finger protein; FXY2 [Homo sapiens]	'human homologue of rat ribosomal protein 1.9' [Homo sapiens]	TIMP [Homo sapiens]	(AF110731)
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2		blastx.2	blastx.2	blacty 2	blastx.2
	1328	1330	1331	1332	1333	1334	1335		1336	1337	1338	1339
	HNOKV82 R	HNORA56 R	HNORB68 R	HNORC14 R	HNORD11	HNORE65	HNORF03R		HNORF04R	HNORF05R	110001120	HNORH03
	HNOKV82	HNORA56	HNORB68	HNORC14	HNORD11	HNORE65	HNORF03		HNORF04	HNORF05	21000101	HNORH03

				B166 [Homo sapiens]				
HNORH05	HNORH05 R	1340	blastx.2	ubiquitin conjugating- protein [Oryctolagus cuniculus]	gb AAA31492.1	100%	43	498
HNORJ08	HNORJ08R	1342	blastx.2	ATPase [Mus	gb AAA88243.1	%06	3	350
	,			musculus		%06	308	496
					-	38%	514	909
HOCMA02	HOCMA02	1344	blastx.2	(AF016365)	gb AAC15862.1	100%	3	497
	RA		,	hexokinase I [Homo		%59	m	497
				sapiens		%98	490	558
·		-	·		-	20%	490	555
HOCMC08	HOCMC08	1347	blastx.2	elongation factor 1	gb AAA18502.1	%68	19	423
	K			alpha [Oryctolagus cuniculus]				
HOCMF27	HOCMF27	1349	blastx.2	sorbitol dehydrogenase	gb AAA66064.1	100%	3	395
-	R			[Homo sapiens]				
337	HOCMG37 HOCMG37	1352	blastx.2	chaperonin (HSP60)	gb AAA36022.1	%86	4	351
	R			[Homo sapiens]		36%	326	424
HOCMI62	HOCMI62R	1353	blastx.2	S19 ribosomal protein	gb AAA89070.1	%19	64	384
				[Homo sapiens]				
HOCMJ47	HOCMJ47R	1354	blastx.2	ribosomal protein L18a	gb AAC18781.1	%86	4	378
				[Homo sapiens]		100%	413	.439
K20	HOCMK20	1355	blastx.2	stathmin [Xenopus	emb CAA50562.1	77%	80	379
	R			laevis]		64%	363	500
907	HOCML06 HOCML06	1356	blastx.2	glyceraldehyde 3-	gb AAA52496.1	%26	123	707
	~			phosphate			_	
				dehydrogenase (EC				•
				1.2.1.12) [Homo				
				sapiens	•		•	

211 330	594 137	200	615	312 416 113	466	511 315	23	485 567 628	706 279 647 273 711 282
2 217	133 24	9	151	97 309 45	80	317 115	388	3 490 566	224 100 495 31 511 139
85% 76%	98% 100%	%88		97% 100% 43%	%66	100%	100%	98% 84% 66%	51% 81% 47% 29% 28% 31%
dbj BAA91374.1	emb CAA55935.1	emb CAA92522.1	gb AAA53505.1	gb AAC25497.1	emb CAA54918.1	emb CAA46716.1	sp Q16465 YZA1_H UMAN	emb CAA68732.1	gb AAC31808.1
(AK000779) unnamed protein product [Homo saniens]	glutathione transferase T1 [Homo sapiens]	plakoglobin [Homo sapiens]	insulin-like growth factor binding protein 5 [Homo sapiens]	(AF028832) Hsp89- alpha-delta-N [Homo	ribosomal protein S15a [Rattus norvegicus]	fau [Homo sapiens]	HYPOTHETICAL PROTEIN (FRAGMENT).	protein phosphatase 2A (AA 1 - 309) [Oryctolagus cuniculus]	(AF059486) putative actin-binding protein DOC6 [Mus musculus]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
1357	1358	1360	1361	1363	1364	1365	1366	1368	1369
HOCML66 R	HOCMM52	HOCMS75	HOCMS87 R	HOCMY53 R	HOCMY61	HOCMY79 HOCMY79	HOCOC38 R	HOCOC94 R	HOCOF35R
HOCML66 HOCML	HOCMIMS	HOCMS75 HOCMS75	HOCMS87	HOCMY53	HOCMY61	HOCMY79	HOCOC38 HOCOC38	HOCOC94	HOCOF35

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460	532	632	561	965			631	301	621	979	620	474	541		412		351	412	455		217			275	408	322	365	39
2	38	396	472	42			2	89	388	441	522	94	482		8			347	396		2	-		m	286	41	225	4
%ZL	46%	25%	43%	% <i>LL</i>			%19	34%	39%	30%	42%	%46	75%		%16		%96	72%	20%		100%			%96	48%	<i>%9L</i>	26%	100%
gb AAA36111.1				gb AAA36582.1			gb AAA60300.1			•		pir S70029 S70029			gb AAA60284.1		emb CAA04940.1	•			emb CAA31412.1			gb AAB54057.1		emb CAA67576.1		
'FKBP52; 52 kD	FK506 binding protein'	[Homo sapiens]	•	Ro ribonucleoprotein	autoantigen (Ro/SS-A)	precursor [Homo sapiens]	spliceosomal protein	[Homo sapiens]				probable	transmembrane protein	TMC - human	ribosomal protein S17	[Homo sapiens]	(AJ001701)	deoxyhypusine	synthase [Homo	sapiens	3-oxoacyl-CoA thiolase	propeptide (424 AA)	[Homo sapiens]	antisecretory factor-1	[Homo sapiens]	collagen (VI) alpha-1	chain [Homo sapiens]	
blastx.2				blastx.2			blastx.2					blastx.2			blastx.2		blastx.2				blastx.2			blastx.2		blastx.2		
1370				1371			1372					1373			1375		1377				1380			1381		1383		
HOCOF50R				HOCO062	ጸ		HOCOP52R				!	HOC0Q13	R		HOCOT70	R	HOCPF69R				HOCPH40R			HOCPH44R		HOCPIS3R		
HOCOF50				HOCO062			HOCOP52					HOC0013	,		HOCOT70		HOCPF69				HOCPH40			HOCPH44		HOCPI53		

136	314 445	563		394	35	427		240	040	536	37	579	176	629	268	ļ	249	-		203	7,2,7	426	512	
23	63 350	<u>е</u>		 80 v	0	<u>м</u>		1	97	39	2	-	- 603	575	98		 		1	m			441	
%98	85%	%88		84%	100%	100%		1,850	%98	100%	91%	%96	48%	40%	%22		65%			100%		%59	24%	
gb AAB48302.1	emb CAA32094.1	gb AAA80153.1		gb AAB65243.1		gb AAA40742.1			emb CAA00898.1	emb CAA27137.1		emb CAA56074.1			gb AAA74903.1		dbj BAA11479.1			dbj BAA75239.1		gb AAA41088.1		
pinin [Bos taurus]	histone H2A.F [Gallus	isoleucyl-tRNA synthetase [Homo	sapiens]	(AF006012)	dishevelled 2 [Homo sapiens]	clathrin-associated	protein 17 [Rattus	norvegicus	TIMP [Homo sapiens]	alpha subunit (aa 1-	394) [Bos taurus]	translation initiation	factor [Homo sapiens]		P311 HUM [Homo	sapiens]	similar to emb-5	protein of C.elegans.	[Homo sapiens]	(AB017018) JKTBP2	[Homo sapiens]	dihydropyridine-	sesitive L-type calcium	channel alpha-2 subunit
blastx.2	blastx.2	blastx.2		blastx.2		blastx.2			blastx.2	blastx.2		blastx.2			blastx.2		blastx.2			blastx.2		blastx.2		
1385	1386	1387		1389		1390	•		1391	1392		1393			1395		1397			1399		1400		-
HOCPM39	HOCPP65R	HOCPP80R	-	HOCPQ59R	,	HOCPQ66R			HOCPO75R	HOCPR01R		HOCPR29R			HOCPR53R		HOCPR77R			HOCPS35R		HOCPU03R		
НОСРМ39	HOCPP65	HOCPP80		HOCPO59		HOCPO66		•	HOCPO75	HOCPR01		HOCPR29			HOCPR53		HOCPR77			HOCPS35		HOCPI 103		

		-		[Rattus norvegicus]				
HOCPU30	HOCPU30R	1401	blastx.2	KDEL receptor [Homo	emb CAA39371.1	%99	342	518
				sapiens]		%69	139	267
						61%	242	334
HOCPU68	HOCPU68R	1402	blastx.2	preprocathepsin B	gb AAA52129.1	%86	m	569
				[Homo sapiens]				-
HOCPV29	HOCPV29R	1403	blastx.2	thrombin inhibitor	emb CAA80373.1	28%	243	593
				[Homo sapiens]				
HOCPV67	HOCPV67R	1404	blastx.2	serine/threonine-protein	gb AAB03269.1	%88	28	327
			•	kinase PRP4m [Mus		87%	345	518
				musculus]	•			
HOCPV72	HOCPV72R	1405	blastx.2	(AF132952) CGI-18	gb AAD27727.1 AF1	91%	33	203
				protein [Homo sapiens]	32952_1	84%	184	318
HOCPW56	HOCPW56 R	1406	blastx.2	ARL3 [Homo sapiens]	gb AAA21654.1	100%	2	502
HOCPW81		1408	blastx.2	cysteine protease Mch2	gb AAC50168.1	91%	7	739
	ద			isoform alpha [Homo				_
				sapiens]				
HOCPX01	HOCPX01R	1409	blastx.2	(AF052514) thymus	gb AAC33563.1	73%	129	440
				specific serine		20%	113	571
				peptidase [Homo	•	87%	999	589
				sapiens]		78%	483	587
HOCPZ76	HOCPZ76R	1411	blastx.2	(AF002705) beta prime	gb AAB88018.1	%66	2	421
•				COP [Rattus	•	77%	423	524
	,			norvegicus]				
HOCQA26	HOCQA26	1413	blastx.2	interferon-inducible	emb CAA59337.1	100%	205	378
	8			protein [Homo sapiens]		71%	410	571
				, ,		47%	360	473
HOCQA37	HOCQA37	1414	blastx.2	lactoyl glutathione	dbj BAA02572.1	100%	136	276

	-	ŀ	Lines (Homo comienc)		85%	63	151
123	1415 blastx.2		proteoglycan core	gb[AAB00774.1]	91%	84	392
		_	protein [Homo sapiens]	mblC 4 437375 11	%66	3	473
	1416 blastx.2		general transcription factor [Homo sapiens]	emological constants			
	1417 blastx.2	┪	Na+-independent	dbj BAA11541.1	92%	4	222
			neutral and basic amino				_
			acid transporter [Homo				
			sapiens		2000	106	410
	1419 blastx.2	2.	calcium binding protein	dbj BAA23325.1	98%	071	111
			Homo sapiens	-11 A A 26167 11	%88	152	637
\sim	1420 blastx.2	73	ribosomal protein L12	gp AAA30137.1			
- 12	_	,	retinoid X recentor	gb AAA42093.1	%08	3	392
1471	DIASIA.2	<u> </u>	alpha (Rattus		26%	424	567
			norvegicus]		,0,0	1	700
1	1423 blastx.2	x.2	(AF095257)	gb AAD03717.1	80%	٦ ٥	175
			heterogeneous nuclear		%60	777	7 7 7
			ribonucleoprotein		%60	1,4	1 0
			C1/C2; haRNP C1/C2		23%	4/4	746
			[Mus musculus]		ì	8	163
Iڪ	1424 blastx.2	×.2	glyceraldehyde-3-	pir JC5370 JC5370	91%	76.	785
			phosphate		24%	1470	9 5
			dehydrogenase (EC		100%	457	49.4
			1.2.1.12), euthermic				
			tissue - desert jerboa			T	424
163	1425 blastx.2	X.2	ribosomal protein L7a	gb AAA60282.1	74%	0	437
1			large subunit [Homo			,	
		:	sapiens				

629	652	242	324				230		274					133	233		167	304	. 158	1.70	164	164	176	164	146	168	161	91
6	254	144	1				3		∞					56	216		9	164	12	12	45	. 39	30	24	36	13	81	53
62%	28%	91%	100%		_		100%		25%					%99	100%		%96	%89	36%	33%	35%	30%	25%	30%	27%	76%	37%	46%
emb CAB06627.1		gb AAC39866.1	gb[AAB32370.1]	- -			emb CAA56072.1		gb[AAC23784.1]					emb CAA01433.1			gb AAF00095.1										;	-
actin [Brugia malayi]		(AF038952) cofactor A protein [Homo sapiens]	prostacyclin-	stimulating factor,	PGI2-stimulating	factor, PSF 1 sapiens]	hnRNPcore protein A1	[Homo sapiens]	(AC004638) amyloid	precursor protein-	binding protein 1	(APP-B1) [Homo	sapiens]	tissue-specific	secretory protein	[synthetic construct]	(AF113514) histone	acetyltransferase	MORF [Homo sapiens]	i i		4						
1426 blastx.2		blastx.2	blastx.2				blastx.2		blastx.2					blastx.2			blastx.2										•	
1426		1427	1428		-		1430		1431					1434			1435										•	
НОСОС94	R	HOCQH81 R	HOCOI31R	,			HOCQM24	R	HODAF78	24				HODCZ52	8	•	HODDI57R		1, 72								٠.	
HOCQG94 HOCQG94		носон81	HOCOI31	,			HOCQM24		HODAF78					HODCZ52			HODDI57											

101	289	207	181	409	153	153	153	153	153	153	153	153	156	153	153	153	153	233	239	239	233	239
9	83	1	7	305	16		-			16		—	_	-	—	7	10	150	156	174	174	153
34%	%99	%86	100%	100%	%82	10%	70%	64%	%99	71%	28%	28%	%09	28%	28%	58%	54%	75%	%29	77%	%06	62%
	gb[AAA91179.1]	gb AAF25683.1 AF2 16754_1	emb CAA70103.1	gb AAA87567.1	gb[AAC50261.1]	-					-											
	Method: conceptual translation supplied by	(AF216754) over- expressed breast tumor	Ran_GTP binding protein 5 [Homo saniens]	selenium donor protein [Homo sapiens]	zinc finger protein	ZNF136 [Homo	saniens]	Carrondes				-										
	blastx.2	blastx.2	blastx.2	blastx.2	blacty 2	7:West									,							
	1440	1449	1453	1459	1460	201									·,							
	HODER57 R	HODFC44R P00	HODF114R	HODF016	TODEOGA	nonroa	4	-												-		
	HODER57	HODFC44	HODFJ14	HODFO16		HODFO04																

329	242	233	236	153	328	328	319	233	337	319	319	314	314	328	319	325	319	325	439	436	453	499	6	493		470	129
159	153	174	153	49	242	269	239	174	242	272	248	159	174	242	251	239	248	239	74	104	7	. 47	7	338		က	31
45%	63%	%08 .	%09	54%	51%	70%	25%	%09	43%	81%	24%	32%	34%	48%	52%	41%	41%	37%	77%	44%	%68	95%	93%	25%		%68	%06
							-									-		-	gb AAA36648.1		gb AAB03694.1	gb AAB32776.1				gb AAB58952.1	splQ9Y6Y5 Q9Y6Y5
																•			pre-mRNA splicing	factor [Homo sapiens]	DNA topoisomerase III [Homo sapiens]	pancreatic	peptidylglycine alpha-	amidating	monooxygenase, 1 1	protein 4 [Homo	IDN4-GGTR14
											,							,	blastx.2		blastx.2	blastx.2				blastx.2	blastx.2
·																			1461		1462	1463				1475	1478
																		-	HODFP51R	,	HODFQ19 R	HODFQ37	<u></u> ۲			HODGB69 R	норензо
																-			HODFP51		норғ019	HODFQ37				НОДСВ69	норднзо

				Maroda				
	R			FROIDIN.	11-315 A A CILL 11	100%	48	116
HODGH43	HODGH43	1479	blastx.2	(AB005878) BYJIS	apjlbAA41013.11	28%	217	363
у НОДОН	HODGH65	1480	blastx.2	IDN4-GGTR14	sp Q9Y6Y5 Q9Y6Y5	%96	41	127
	R			PROTEIN.	114 A DECOAT 11 A E1	30%	95	475
HODGN53	HODGN53	1482	blastx.2	(AF186461) ring tinger	86461 1	2//	?)
	*			norvegicus]			1	
HODGO52	HODGO52	1487	blastx.2	unnamed protein	emb CAB42187.1	85%		48
20X00011	R			product [unidentified]		75%	110	107
HODGW9	HODGW9 HODGW91	1489	blastx.2	phosphofructokinase	gb AAA79220.1	100%	394	347
1	7	999	0 1	Trough Capture	obl A A 28195.11	49%	190	462
HODGZ63	HODGZ63	1493	blastx.2	putative [Caenorhabditis		21%	470	532
				elegans		ì	č	5
норнр23	норнр23	1495	blastx.2	fused-ccdB	emb CAA71575.1	%76	†	101
	~			Eschericina con	11. 0.1001	10007	24	323
HODHE54	HODHE54	1498	blastx.2	lysyl oxidase-like	gb[AAA50162.1]	0,001	323	418
	&			protein [Homo sapiens]		32%	94	348
HODHE88	HODHE88	1499	blastx.2	(AK001737) unnamed	dbj BAA91871.1	83%	29	196
	ద			protein product [Homo				
HODHGS6	HODHG56	1500	blastx.2	(AF151888) CGI-130	gb AAD34125.1 AF1	100%	121	231
Z 2	2		_	protein [Homo sapiens]	51888 1	,000	(200
HODHK 86	норнк86	1503	blastx.2	(AB005299) BAI 3	dbj BAA25363.1	93%	7 7	707
	R R			[Homo sapiens]		48%	207	590
						2004	730	126
35 11 00 11	HODII.36R	1505	blastx.2	(AL031177)	emb[CAA20118.1]	40%	477	202
טכעונעטט	11010101011							

118		558	338	473	333		424	286	391	207	464	624		344	651	682	538	•			424		489	512	
26		. 64	3	348	503		65	41	38	463	417	1		3	331	416	5				116		16	450	
32%		%86	%96	28%	%99		%66	45%	34%	100%	75%	%66		%96	71%	28%	%9/				%85		%08	47%	
		emb CAA66039.1	emb CAB06294.1		emb CAB51405.1		dbj BAA08302.1		-	٠	•	gb AAA60289.1		gb AAA36470.1	-		emb CAA45089.1				gb AAF32294.1 AF2	16306 1	gb AAD22669.1 AC0	07193_3	
dJ889N15.2.2 (26S	Proteasome subunit p28 (Ankyrin 1	prepronociceptin [Homo sapiens]	serine/threonine kinase	[Rattus norvegicus]	(AL096881)	hypothetical protein [Homo sapiens]	calmodulin [Homo	sapiens]				ribosomal protein S6	[Homo sapiens]	acidic ribosomal	phosphoprotein (P0)	[Homo sapiens]	homologue to	elongation factor 1-	gamma from A.salina	[Homo sapiens]	(AF216306) DCRC-1	[Mus musculus]	(AC007193)	PPP5_HUMAN [Homo	sapiens
		blastx.2	blastx.2		blastx.2		blastx.2					blastx.2		blastx.2			blastx.2				blastx.2	•	blastx.2		
	,	1506	1508		1514		1519					1520		1521	•		1526				1527		1528		
А		HODJZ09R	HODKB82	R	HODKD64	R	HODKK26	ጸ				HODKK40 HODKK40	R	HODKK73	8		HODKN65	&			HOECR39R		HOFAB77R	-	
		HODJZ09	HODKB82		норкр64		HODKK26					HODKK40		HODKK73			HODKN65	·		ŕ	HOECR39		HOFAB77		

HOFMF79	HOFMF79 HOFMF79	1530	1530 blastx.2	cytochrome b [Canis	gb AAD04775.1	%08	4	339
	₹			[familiaris]		38%	305	430
HOFMJ88	HOFMJ88R	1531	blastx.2	receptor protein-	gb AAA61139.1	%9L	17	370
				tyrosine kinase [Homo sapiens]	•			
HOFMM8	HOFMM84	1533	blastx.2	hydroxymethylglutaryl-	pir A43533 A43533	72%	112	252
4	ĸ			CoA reductase		85%	254	274
				(NADPH) (EC				
				(fragments)	,			:
HOFMN93	HOFMN93	1534	blastx.2	(AF133669) ARL-6	gb AAD33046.1 AF1	61%	36	299
	N N			interacting protein-1	33669_1	84%	214	327
				[Mus musculus]				
HOFMP59	HOFMP59	1535	blastx.2	mitochondrial	dbj BAA01387.1	<i>%LL</i>	П	360
	٠ د			acetoacetyl-CoA		83%	369	422
				thiolase [Homo				
				sapiens]				
HOFMT68	HOFMT68	1536	blastx.2	RNAse L inhibitor	gb AAC24730.1	25%	137	547
	R			[Mus musculus]		40%	264	581
HOFMT69	HOFMT69	1537	blastx.2	HETEROGENEOUS	sp P52272 ROM_HU	%08	127	264
	2			NUCLEAR	MAN	75%	27	122
				RIBONUCLEOPROTE		36%	86	316
				IN M (HNRNP M).				
HOFMU92	HOFMU92 HOFMU92	1538	blastx.2	(AP000694) chromatin	dbj BAA89426.1	63%	124	432
	<u>س</u>			assembly factor 1,			,	
				subunit B (p60) [Homo				
				sapiens]				
HOFNF63	HOFNF63R	1540	blastx.2	phopshomannomutase [Homo sapiens]	gb AAC51368.1	70%	76	478
HOFNF76	HOFNF76R	1541	blastx.2	(AB002806) OS-9	dbj BAA24363.1	94%	17	370

	414	186	232	009	397	345	242 460 270	219 219 115 350
	157	19	17	16	80	22	18 245 244	115 109 35 204
	%62	%96	91%	74%	84%	%86	98% 62% 100%	85% 57% 64% 36%
	dbj BAA21881.1	gb AAA30678.1	emb CAB52550.1	gb AAA60055.1	sp Q29386 Q29386	gb AAB02696.1	gb AAD17375.1	gb AAA39906.1
isoform 2 is missing nt 1642-1806; OS-9 1	(AB006679) ATP binding protein [Homo sapiens]	osteonectin precursor [Bos taurus]	(AL034417) bK215D11.1 (RNA-binding protein regulatory subunit) [Homo sapiens]	pyruvate dehydrogenase E1- alpha precursor [Homo sapiens]	KERATIN TYPE II CYTOSKELETAL 8 (FRAGMENT).	activin receptor type I [Bos taurus]	(AF129075) T- COMPLEX PROTEIN 1, THETA SUBUNIT (TCP-1-THETA) [Homo sapiens]	protein disulfide isomerase [Mus musculus]
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	1542	1543	1544	1545	1546	1547	1548	1549
	HOFNG51 R	HOFNK44 R	HOFNY53 R	HOFOB65R	HOFOB79R P00A	HOFOE22R	HOFOF47R	HOFOF56R
	HOFNG51	HOFNK44	HOFNY53	HOFOB65	ноғов79	HOFOE22	HOFOF47	HOFOFS6

HOGCF48					_	40%	-	777
348						7027	245	592
	HOGCF48R	1553	blastx.2	hepatitis C-associated	dbj BAA06043.1	%96	92	240
				microtubular aggregate		67%	414	587
				protein p44 [nound		75%	523	570
TOGOGR3	HOGCG83	1554	blastx.2	keratin [Homo sapiens]	emb CAA31695.1	79%	309	310
	R					26%	21	113
		_		1 mom	gb AAA85332.1	93%	3	179
HOGCI28	HOGCI28R	1556	blastx.2	component C3 [Homo		83%	173	337
			!	sapiens		7000	103	471
HOGCJ10	HOGCJ10R	1557	blastx.2	(AF032667) rexo70	gb AAC015/9.1		}	
: :				Rattus norvegicus	11 A MT 59K1 11	%56	13	459
11000165	HOGOISSR	1559	blastx.2	Grb14 [Homo sapiens]	gb AAC13001.1	10007	31	183
TOGCT44	HOGCT44	1562	blastx.2	casein kinase I-alpha	gb AAC41/60.1	2001	<u> </u>	}
-	2			[Homo sapiens]	11 4 4 4 6 9 6 2 6 11	%92	3	521
HOGCV85	HOGCV85	1563	blastx.2	heparin binding protein	golada a social	73%	368	535
				Homo sapiens	11 005017 11	70%	43	504
HOGCV93	HOGCV93	1564	blastx.2	elongation factor 1	gb AAA18302.1	2	<u> </u>	·
	K			alpha [Oryctolagus				- 1
				cumounds	phAA418502.1	97%	∞	340
HOGCY12		1565	blastx.2	alpha [Oryctolagus		41%	302	547
	ፈ			cuniculus		0,40%	6	329
HOGCY58	HOGCY58	1566	blastx.2	lipocortin II [Homo	dbj BAA00013.1 	76%	332	517
) i l				sapiens		30%	150	326
		1567	blocky 2	extracellular matrix	emb CAA68724.1	100%	m	260
HOGCY74	HOGCY /4	1001		protein BM-40 (AA 1 -				

				303) [Homo sapiens]				
HOGDD29	HOGDD29	1568	blastx.2	drebrin E2 [Homo	gb AAA16256.1	%88	e.	152
				sapiens				
НОСРС03	НОСРС03	1570	blastx.2	I-caldesmon II [Homo	dbj BAA14419.1	%56	191	388
	ጸ			sapiens]		87%	393	491
HOGDI44	HOGDI44R	1572	blastx.2	neutrophil gelatinase	emb CAA58127.1	100%	55	192
	•			associated lipocalin				
HOGD025	HOGDO25	1574	blastx.2	XIAP associated factor-	emb CAA68030.1	100%	2	163
	R			1 (ZAP-1) [Homo	-			·····
				sapiens]				
HOGDO58	нодрозя	1575	blastx.2	acidic ribosomal	gb AAA36470.1	%59	98	373
	R			phosphoprotein (P0)				
				Homo sapiens	•			
HOGDP10	HOGDP10 R	1576	blastx.2	keratin [Homo sapiens]	emb CAA73943.1	%68	m	287
HOGD095	нодроэ5	1578	blastx.2	FUSE binding protein 3	gb AAC50893.1	%06	20	435
	X			[Homo sapiens]	•	32%	157	378
	:			1	-	62%	390	461
						48%	413	493
HOGDR70	HOGDR70	1579	blastx.2	gamma subunit of CCT	emb CAA52808.1	83%	102	278
	&	•		chaperonin [Homo sapiens]				
HOGDV93	HOGDV93	1580	blastx.2	glyceraldehyde 3-	gb AAA52496.1	%86	4	414
				phosphate	-			
				dehydrogenase (EC				-
	•			1.2.1.12) [Homo				
				sapiens]				
HOGEA27	HOGEA27	1581	blastx.2	cytokeratin 8 [Homo	gb AAA35763.1	81%	2	160
			:		-			

	2			sapiens	11 07700 11	75%	3	482
HOGED85	HOGED85 R	1582	blastx.2	AP-3 complex beta3A subunit [Homo sapiens]	gb AAD03778:1	85%	305	445 173
	SCAGOOT	1583	hlactx 7	HLA-Aw34.2 antigen	emb CAA43874.1	91%	38	316
HOGEK22	HOGENES	5		Homo sapiens]		020%	140	322
HOGEN30	HOGEN30	1584	blastx.2	regulatory subunit RI	emb CAA01027.1	0	2	
	24			alpha Homo sapiens	11 4 A C 22 12 K 11	75%	83	205
HOGEP69	HOGEP69R	1587	blastx.2	(AF056182) G-protein	gb AAC33430.1	78%	219	260
				remericella nidulans				656
	TIO OTTOOT	1500	blacty 2	Plakoglobin [Homo	gb AAA64895.1	%49	CCI	700
HOGE1.60	HOGETON			saniens			1	7:7
110001/68	HOGEW58	1590	blastx.2	focal adhesion kinase	gb AAA35819.1	%96	4	4 14 ·
HOGE W 28	D			[Homo sapiens]		7020	53	144
HOGE703	HOGEZ03R	1592	blastx.2	kinase [Homo sapiens]	emb CAA65450.1	83%	187	53.
					1 100000	7900	373	543
STOOTH 69		1601	blastx.2	ZZ:beta-Gal' IgG-	gb AAB00807.1	3270	<u> </u>)
HOOHEOS	R -	<u> </u>		binding fusion protein				
				unidentified cloning 1	- 5000	7000	-	156
HOOHP84	HOOHP84	1602	blastx.2	(AF013215) ribosomal	gb[AAB65437.1]	0/70	4	
	_		-	protein 32 [DOS tautus]	ohlAAA72865.1	%69	292	423
нооноон	ноонооб	1603	blastx.2	pixolue lusion process				000
110011002	$\overline{}$	1604	blastx.2	(AB000910) ribosomal	dbj BAA19210.1	100%	ç 8	745
מאשטטש		_		protein [Sus scrofa]		2000	326	458
TOOTE	_ i	1605	blastx.2	pfxblue fusion protein	gb AAA72865.1	73%	350	505
TONE OF I				[synthetic construct]		0/0/	1	100
2100110011	· TIOOUT13	1608	hlastx.2	steroid receptor TR2-11	gb AAA36761.1	95%		201
HOOHITS	_		-					

	R			[Homo sapiens]		%56	193	264
HOOIA46	HOOIA46R	1610	blastx.2	cpn10 protein [Bos taurus]	emb CAA49288.1	100%	3	239
HOOIB54	HOOIB54R	1613	blastx.2	ribosomal protein L23a [Homo sapiens]	gb AAB17510.1	75%	68	199
HOOIG71	HOOIG71R	1615	blastx.2	ribosomal protein L27 [Homo sapiens]	gb AAA19815.1	%86	2	250
HOOJN84	HOOJN84R	1624	blastx.2	IEF 7442 [Homo sapiens]	emb CAA51360.1	100%	4	189
HOOJT32	HOOJT32R	1629	blastx.2	ribosomal protein L23a [Homo sapiens]	gb AAA03341.1	%9 <i>L</i>	34	237
НООЈТ65	HOOJT65R	1630	blastx.2	(AJ388520) Ribosomal protein [Canis familiaris]	emb CAB46822.1	%88	43	123
H00JY44	HOOJY44R	1636	blastx.2	human homologue of rat phosphatidylethanolami ne binding protein [Homo sapiens]	dbj BAA03684.1	100%	2	1.84
ноолу80	HOOJY80R	1637	blastx.2	cytochrome b5 [Homo sapiens]	gb AAA35729.1	%86	61	462
ноолу92	HOOJY92R	1638	blastx.2	mitochondrial matrix protein [Homo sapiens]	gb AAA60127.1	98% 91% 66% 62%	48 403 513 341	344 513 686 445
HOOKN43	HOOKN43 R	1640	blastx.2	ZZ:beta-Gal' IgG- binding fusion protein [unidentified cloning 1	gb AAB00807.1	%89	124	399
HOPJF95	HOPJF95R	1643	blastx.2	nucleolar protein p40	gb[AAB46731.1]	91%	209	511

219	461	280	551	594	476	460	543	49	655		432	017	212	000				594	709	653	380	417					460	
130	237	98	m	535	42	182	172	2	7			77	בי ל ל	670		_		34	584	615	09	379					7	
% 06 .	%58	100%	%16	27%	46%	44%	34%	100%	%56			7000	9,000	83%				%66	%05	84%	34%	61%		•			100%	
	gb AAF28946.1 AF1	61386 1	emb CAA42052.1		emb CAA27268.1				pir A42008 A42008		gb AAA36597.1	11 000000000000000000000000000000000000	emb CAB/3309.1					emb CAA43019.1			gb AAC26291.1			-			gb AAA51690.1	
[Homo sapiens]	3PC268		protein	[Homo sapiens]	C gamma 3 [Homo	saniens	- Company		homeotic protein	HOX4E - human	scar protein [Homo	Sapicins	(AL050318) dJ977B1.5	(myosin regulatory	light chain 2, smooth	muscle isoform) [Homo	sapiens]	elongation factor-1-	beta [Homo sapiens]	•	(AF077539) contains	similarity to human	melanoma antigen p15	(GB: U19796)	[Caenorhabditis	elegans]	aldolase A (EC	4.1.3.13) [Homo sapiens]
	blastx.2		blastx.2		blastx.2				blastx.2	•	blastx.2		blastx.2					blastx.2			blastx.2						blastx.2	
	1644		1646		1650	2			1656		1659		1660					1661			1662						1663	
1	HOPJG60R))))	HOPIG79R) ; ;) ;	нортибя	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,			HOPKA06	~	HOPKG16	소	HOPKG47	~				HOPKG83	2	ı	HOPKK38	R					HOPKN14	~
	HOPIG60		HOPIG70	2001	уундон	COTTE TOTT			HOPKA06		HOPKG16		HOPKG47					HOPKG83			HOPKK38						HOPKN14	

483	81	405	527			435	435	498		173	514	155	155	155	170	152	203	155	- 221
31	1	4	285	154	494	385	115	28		9	5	12	21	21	21	21	21	,	77
94%	%96	%92	%06	%06	41%	35%	%78	%68	•	100%	%98	85%	%08	75%	70%	79%	%09	1000	0/.0/
emb CAA38722.1	emb CAB70733.1	dbj BAA11824.1	gb[AAB34216.1]				gb[AAA60286.1]	gb AAC16388.1		emb CAA02873.1	gb[AAC15856.1]	emb CAB45723.1		•					
17,000 dalton myosin light chain [Bos taurus]	(AL137423) hypothetical protein Homo saniens1	Six5 [Mus musculus]	PAX8=paired-box-	protein [alternatively	spliced] [human, 1		ribosomal protein S20 [Homo sapiens]	(AF063243) ribosomal protein L30 [Bos	taurus	unnamed protein product [Homo sapiens]	ribosomal protein L11 [Homo sapiens]	(AL080125)	hypothetical protein	[Homo sapiens]					
blastx.2	blastx.2	blastx.2	blastx.2	_	-	,	blastx.2	blastx.2		blastx.2	blastx.2	blastx.2							
1664	1665	1666	1667				1668	1669		1670	1671	1674		,				•	
	HOPKO04 RA	HOPKO61 R	HOPKP45R	-			HOPKQ20 R	HOPKQ82 R		HOPKR56R	HOPKU33 R	HOVBK49	X				-		
HOPKN67	HOPKO04	HOPKO61 HOPKO61	HOPKP45				НОРКQ20	НОРКО82		HOPKR56	НОРКИЗЗ	HOVBK49				-			

152	152	369	375	375	372	375	170	375	375	375	173	407	369	407	407	407	407	407	407	401	407	152	305	146	434	109	249	319	399
12	21	112	121	121	121	121	24	121	121	121	27	258	178	258	264	261	264	258	258	261	258	27	243	42	246	20	112	182	340
61%	63%	40%	40%	38%	38%	36%	46%	34%	34%	32%	38%	42%	34%	36%	37%	34%	39%	34%	32%	31%	30%	36%	25%	78%	%05	25%	34%	%98	75%
	,									-			-				,					-			gb AAD46422.1 AF1	05715 1	•	gb AAB81618.1	
																							***		(AF105715) ubiquitous	miclear profein [Gallus	nuclear protein [current	(AE02531) recentor	for activated protein
	-						_		-			-			1				-						blocky 2	Ulasta:4		1-10mm 2	OldSLX.2
-					_	-	_								-			_							1675	7/01		1000	0/01
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	•	_			····							-			-							-			0 0 0	HOVBS15			HOVBX41

341	214	578	321 454	220 464	522 196 190	361 506	204	282	226 98 210	299
309	23	36	287	372	463	444	91	112	116	15 405
72%	%99	72%	92%	97%	23%	65% 61%	%68	%96	75% 81% 27%	98%
	emb CAA06702.1	pir S03957 R5RT18	emb CAA43139.1	emb CAB70827.1		pir A44282 A44282	gb AAD32925.1 AF0 67728_1	gb AAB00807.1	emb CAA40254.1	gb AAA36032.1
kinase C [Oreochromis niloticus]	(AJ005798) thyroid hormone receptor alpha 2 [Sus scrofa]	ribosomal protein L18a - rat	alpha-smooth muscle actin [Oryctolagus cuniculus]	(AL137585) hypothetical protein	[Homo sapiens]	retrovirus-related pol polyprotein pseudogene - human 1	(AF067728) transactivating protein BRIDGE [Rattus norvegicus]	ZZ:beta-Gal' 1gG- binding fusion protein [unidentified cloning 1	23 kD highly basic protein [Homo sapiens]	binding protein [Homo sapiens]
•	blastx.2	blastx.2	blastx.2	blastx.2		- blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	1678	1680	1681	1682		1684	1686	1688	1692	1693
	HOVCN03 R	HOVCU89 R	HOVCV33 R	HOVCW45 R		HOVCZ45 R	HOVDB61 R	HOVDB65 R	HOVDG71 R	HOVDH09 R
·	HOVCN03	68UDVOH	НОУСУЗЗ	HOVCW4 5		HOVCZ45	ноурв61	ноурв65	HOVDG71	ноурно9

381	707		254	611	612	393		200	47	369			576	2		186	}		648		717	\perp	221	
316	121	171	126	370	240	49		90	21	277	·		ac.	07		4	•		64	3	ડ -	3	217	
81%	44%	81%	%69		~1% 80%	%66		è	77%	03%) 600 F	\$001 ·		1000%	2001		92%		82%			
		gb AAB00807.1	CHIA A D34103.11AF1	51866 1	gb AAA67526.1	LL A A A 73877 11	li.,,,oc,,vvvvlag		gb[AAA52135.1]	11 200000 11 111	gb AAB00807.1			emb CAA58449.1			gb AAA21814.1		oh AAA36161.1	0.0	emb CAA00596.1		emb CAA40254.1	
			1	(AF151866) CGI-106	75 [Homo	sapiens	thyroid receptor	sapiens	cytochrome c-1 [Homo	sapiens	ZZ:beta-Gal' IgG-	binding fusion protein	funidentified cloning 1	Lutheran blood group	glycoprotein [Homo	sapiens	macrophage migration	inhibitory factor [Homo	sapiens]	Iaminin-binding protect	LT A DR. heta. A	[Homo saniens]	23 kD highly basic	protein [Homo sapiens]
		blastx.2		blastx.2	blastx.2		blastx.2		blastx.2		blastx.2			hlastx.2			blastx.2		١	blastx.2		blastx.2	blastx.2	
		1694	_	1696	1703		1704		1707		1712	1		1717	(1)1		1718			1719		1720	1721	
	-	HOVDH75		HOVDV70			HOVEF81R		HOVELSIR		HOVEVS8	2017	4	TTOX11100D	WOOM AND A		TOVIE SER	NOCYTE OUT		HOVJU75R		HOVJW17	R HOVIV68R	
		HOVDH75		HOVDV70	HOVEF34		HOVEF81		מסעיפו לו	TOTAL A CILI	_	HOVEIO			HOVIJON		23 0115011	HOVJKO		HOVJU75		HOVJW17	89/11/1011	2017/04

241	487	379	601	629	929	553	684	253	299	331		929			`	786	805		464		456			451	
149	2	2	353	2	65	65	496	17	231	140		68				64	746		84		136			56	
35%	%86	%86	%99	%08	41%	31%	38%	74%	95%	%05		%46				%88	20%		%26		%66			%08	
	emb CAB69339.1	emb CAA27801.1	gb AAA36776.1	gb[AAD08718.1]				gb AAA87172.1		gb AAA57126.1		emb CAA56869.1		-		emb CAA40268.1	-		dbj BAA04547.1		gb AAA60602.1			emb CAA47406.1	
	unnamed protein product [unidentified]	PBG-D (aa 1-344) [Homo sapiens]	transformation-related protein [Homo sapiens]	poly(A)-binding	protein [Homo sapiens]		•	heat-shock protein	[Canis familiaris]	ADP-ribosylation	factor [Homo sapiens]	Nascent polypeptide	associated complex	alpha subunit [Homo	sapiens]	protein synthesis	initiation factor 4A	[Mus musculus]	HBp15/L22 [Sus	scrofa]	isolog of yeast suil and	rice gos2; putative	[Homo sapiens]	amphiglycan [Homo	sapiens
	blastx.2	blastx.2	blastx.2	blastx.2				blastx.2		blastx.2		blastx.2			,	blastx.2			blastx.2		blastx.2			blastx.2	
	1723	1725	1726	1727				1729		1730		1731				1732			1733		1734			1735	
	HOVKE66 R	HOVKG18 R	HPAMB11 R	HPAMB60	R			HPAMB93	R	HPAMC04	ጸ	HPAMC19	~	•		HPAMC27	<u>~</u>		HPAMC90	R	HPAMD56	~		HPAME35	¥
	HOVKE66	HOVKG18	HPAMB11 HPAMB11	HPAMB60				HPAMB93		HPAMC04		HPAMC19 HPAMC19				HPAMC27 HPAMC27			HPAMC90		HPAMD56 HPAMD56			HPAME35 HPAME35	

414 520	434		612	374	380	516	661	701	592		397	609	494	190		209				493	009	354	358	159	
178	63	3	331	117	7,	304	518	627	11		65	412	405	7		381				2	472	4	221	46	
45% 73%	47%	200	61%	%65	7070	%4% %50	72%	%89	87%		%1.6	64%	989	100%		100%				%86	%0%	7009	91%	42%	
gb AAA74186.1	114 4 100004 114 11	gb AAD2//45.1 AF1 32970_1	gb[AAC18792.1]			gb[AAB96924.1]			emblCAA47670.1		emb CAA81022.1	•		emblCA A40254.11		gb AAC36129.1					emolCAS+CAS-1.1	11 7 00000 11	gb AAC9/980.1		
actin [Dictyostelium discoideum]	7	(AF132970) CGI-36	1	117	gb X62724 from 1 1	(AF039752) histone	deacetylase-2; HD-2	Gailus gailus]	ribosomal protein S8	[Homo sapiens]	rihosomal protein S7	Homo conjenel	[choides onion]	02 1-D Lichly, bosio	23 KD mgmy basic protein [Homo saniens]	(AF017153) putative	RNA helicase and	RNA dependent	ATPase [Mus	musculus	23 kD highly basic	protein [Homo sapiens]	(AC006127) BRG-1-	HUMAN AAA 812-	GRB1; 1 1 SNF2L4
blastx.2		blastx.2	bloch 2	Ulasta:		blastx.2			Llower 2	DIASLX.2	blacty 2	Uldath:			blastx.2	blacty 2	- Creation				blastx.2		blastx.2		•
1737		1738	┰			1740			-1-	1/41	17/12	74/1			1744	1775	£/1				1746		1748		
9	4	HPAMF38	K	HPAMG44		HPAMG54	X		4, 14, 14, 14, 14, 14, 14, 14, 14, 14, 1	HPAMIIK	TIPANATAID	HFAIMJ/IR			HPAMQ76	K TIDAN (TEAT)	HL AIM I	4			HPAMU33	8	HPAMW44	꿈	
HPAMF16 HPAMF1		HPAMF38		HPAMG44		HPAMG54			- 1	HPAMIII	-	HPAMJ/1			HPAMQ76		HPAM14/				HPAMU33		HPAMW4	4	

	563 297	508 600	149	599 676	487 729 605	678 678 697	376	235	412	. 624
	294	20 508	3	581	107 466 579	7 19 626	.2	'n	∞	13
	95% 92%	93% 61%	%56	99% 95%	99% 61% 88%	92% 34% 58%	100%	%86	%66	%56
	gb AAD05196.1	gb AAA42278.1	emb CAB44357.1	emb CAA32847.1	gb AAA36033.1	gb AAA59954.1	gb AAA36033.1	gb AAC27653.2	gb AAA60284.1	emb CAA40254.1
[Homo sapiens]	(AC004908) similar to ribosomal protein L.23a; similar to P.29316 (PID:g132848) [Homo sapiens]	triosephosphate isomerase [Rattus norvegicus]	(AJ006776) IF2 protein [Homo sapiens]	SP-40,40 prepropetide (AA -22 to 427) [Homo sapiens]	cell surface glycoprotein [Homo sapiens]	nucleolin [Homo sapiens]	cell surface glycoprotein [Homo sapiens]	(AF078820) high mobility group protein [Spalax ehrenbergi]	ribosomal protein S17 [Homo sapiens]	23 kD highly basic
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	1749	1750	1751	1752	1753	1754	1755	1757	1759	1760
	HPAMY45 R	HPAMZ14 R	HPAMZ15 R	HPAMZ81 R	HPANA05 R	HPANA07 R	HPANA28 R	HPANB32R	HPANE49R	HPANE52R
	HPAMY45	HPAMZ14	HPAMZ15	HPAMZ81	HPANA05	HPANA07	HPANA28	HPANB32	HPANE49	HPANE52

	592 174	119	090						179		335		7			427		306			414			356	314		
	119 34	3	-	8					36		99		337			53	1	136			244			54	~	, ·	
	82% 48%	100%	ì	3.7%					100%		%86	<u>·</u>	100%	200		7088	0/00	94%			%70	2		94%	7000	3//6	
	emb CAA32033.1	dbj BAA00013.1		emb CAB01127.1	•				110 A A 2655A 11	emolecaecoloma	1.1 A A CO300 11	gplagaouzou	TT I V ZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZ	sp[Q16465] Y ZA1_H	UMAIN	100000000000000000000000000000000000000	gb AAA16105.1	11 200000 11	gp AAB0080/.1		11 200000 11	gb AAB0080/.1		114 4 4 8 6 4 4 6 2 1 1	gb AAAoo405.1	emb CAAZ/13/.1	
I Trans caniens	nase	B [Homo sapiens]		d using	nilarity	to Drosophila RNA 1	this gene	[Caenorhabditis	\dagger	IZA.Z (AA 1-	127) [Bos taurus]	ribosomal protein L37a	Homo sapiens	HYPOTHETICAL	PROTEIN	(FRAGMENT).	ribosomal protein	[Homo sapiens]	ZZ:beta-Gal' IgG-	binding fusion protein	[unidentified cloning l	ZZ:beta-Gal' IgG-	binding fusion protein	[unidentified cloning 1	Csa-19 [Homo sapiens]	alpha subunit (aa 1-	394) [BOS tam us]
	blastx.2	Cathol	DIASUX.2	Llooter 2	7.0001/					blastx.2		blastx.2		blastx.2			blastx.2		blastx.2			blastx.2			blastx.2		
	1761 b		1/62		00/1					1770		1771	:	1772			1773		1778			1780			1781	1784	
	HPANE87R 1		HPANJ67R 1	十	HPCOJS9K			· 		HPCOL81R		HPCO090R		HPCOO95R			HPCOP23R		HPCORSOR	111		HPCOV35R	ALCCY OO HI		TECOXIA1B	HPCPD26R	
	HPANE87 H	\neg	HPANJ67 H	7	HPCOJ59 F					TIPCOT 81		T 06000dH		╅	Trecoon		11000003		十	HECONT		3000001	HPCOV33			HPCPD26	

101	185	327 451	326 40	909	190	544	379	177	306
3	3	422	222 2	336	2	374	209	58	1
100%	%19	%06 %E6	97% 92%	% 96	%86	%96	%96	%76	%66
dbj BAA23593.1	pir JC2022 JC2022	emb CAA80977.1	gb AAA60282.1	gb AAB00807.1	emb CAA73248.1	gb AAB00807.1	gb[AAB00807.1]	.gb AAB28826.1	emb CAA42782.1
antizyme inhibitor [Homo sapiens]	enhancer factor I chain A-D - rat	integrin associated protein [Homo sapiens]	ribosomal protein L7a large subunit [Homo sapiens]	ZZ:beta-Gal' IgG- binding fusion protein [unidentified cloning 1	putative progesterone binding protein [Homo sapiens]	ZZ:beta-Gal' IgG- binding fusion protein [unidentified cloning 1	ZZ:beta-Gal' IgG- binding fusion protein [unidentified cloning 1	major histocompatibility complex class II antigen beta chain, 1 Partial Mutant, 57 aa] [Homo sapiens]	H+-ATP synthase subunit b [Homo sapiens]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
1788	1790	1797	1800	1802	1803	1804	1806	1807	1809
HPCPH52R	HPCPU27R	HPCQT88R	HPCQX47R	HPCTD21R	HPCTD23R	HPCTD25R	HPCTF29R	HPCTF83R	HPCT186R
нРСРН52	HPCPU27	HPCQT88	HPCQX47	HPCTD21	HPCTD23	HPCTD25	HPCTF29	HPCTF83	HPCTI86

538	377	148 248 318	205	153	254 342	412	542	209 85 244
77	84	26 138 250	20	1	3 229	7	45	84 14 209
%66	%66	85% 91% 100%	85%	100%	95% 64%	100%	84%	95% 54% 91%
gb AAC05826.1	emb CAA27193.1	emb CAA56100.1	emb CAB55934.1	dbj BAA08389.1	gb AAA67551.1	emb CAA87392.1	gb AAA59831.1	emb CAA50506.1
(AF042857) lung cancer antigen NY-LU-12 variant A [Homo sapiens]	ribosomal protein L35a (aa 1-110) [Rattus norvegicus]	8.2 kDa differentiation factor [Homo sapiens]	(AL117452) hypothetical protein [Homo sapiens]	Aopl_Human, MER5(Aopl_Mouse)- like protein [Homo sapiens]	2,4-dienoyl-CoA reductase [Homo sapiens]	RNA polymerase II elongation factor-like protein [Homo sapiens]	MHC HLA-DR-beta chain precursor old gene name 'HLA-DRA-DRA1' [Homo sapiens]	40S ribosomal protein S14 [Podocoryne carnea]
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
1811	1815	1816	1817	1818	1822	1823	1824	1825
HPCTO69R	HPCTV40R	HPCTV53R	HPCTV92R	HPCTX22R	HPDOF81R	HPDOP05R	HPDOP20R	HPDOS87R
HPCTO69	HPCTV40	HPCTV53	HPCTV92	HPCTX22	HPDOF81	HPDOP05	НРДОР20	HPDOS87

99	269	468		489	485			291	110	519		460			214		337	319	319	319	399	402	387	384	402
. 1	3	250		64	318			121	87	349		131			2		128	125	125	155	292	325	325	325	325
54%	%86 ·	%68		94%	%86			92%	87%	25%		94%			<i>%9L</i>		45%	36%	36%	32%	37%	48%	27%	%09	42%
	gb AAA51828.1			gb AAA63263.1	gb AAC63516.1			gb AAA36021.1		gb AAB00807.1		emb CAA08974.1			gb[AAC78797.1]		dbj BAA86989.1								-
	N-acetyl-beta-	glucosaminidase	prepro-polypeptide [Homo sapiens]	tripeptidyl peptidase II [Homo sapiens]	(AF073298) small	EDRK-rich factor 2	Homo sapiens	Q1Z 7F5 [Homo	sapiens]	ZZ:beta-Gal' IgG-	binding fusion protein funidentified cloning 1	(AJ010046) guanine	nucleotide-exchange	factor [Homo sapiens]	(AF053356) ORF3,	splicevariantc [Homo sapiens]	(AB021643)	gonadotropin inducible	transcription repressor-	3 [Homo sapiens]					•
	blastx.2			blastx.2	blastx.2			blastx.2		blastx.2		blastx.2			blastx.2		blastx.2					•		•	
	1826	•		1827	1829			1830		1835		1836			1837	· · · · ·	1838			·					
	HPDOUS4	~		HPDOZ43R	HPDPC90R	V		HPDPH14R		HPDPQ16R		HPDPR73R		-	HPDPS51R		HPDPS90R		•			•		•	
	HPDOU54			HPDOZ43	HPDPC90			HPDPH14		HPDPQ16		HPDPR73			HPDPS51		HPDPS90								

518	2	407	461	437	418	393	387	642	642	453	642	689	692	692	334	271	494	267	240	539				441			
390		275	399	396	14	199	199	7	7	7	226	639	639	639	89	r	345	226	178	18				-			
32%		38%	47%	64%	100%	%96	85%	%56	94%	97%	94%	64%	61%	61%	. %001	2000	94%	85%	41%	%66				%96		•	T.
					gb AAA60284.1	oblAAA60213.11		eb[AAB92373.1]		•					gb AAA60280.1		gb AAD09822.1			gb AAA20683.1				emb CAA18608.1			
					ribosomal protein S17	arothimosin alpha	Homo saniens	(AF038129)	polynipianitin fovis	port acream Loves	[corns			-	ribosomal protein L37a	Homo sapiens	(AF115850) PAR	protein automore		succinate	dehydrogenase	flavoprotein subunit	[Homo sapiens]	(AL022577) dJ353H6.2.2	(SW1/SNF related,	matrix associated, 1	(SINFALLI)
					blastx.2	blocks 2	Oldsta.2	Flacty 2	Oldstar						blastx.2		blastx.2			blastx.2				blastx.2			
					1841	1040	7401	10/3	Crot				:	_	1845		1846			1847				1848			
					HPDPX12R	FUUD	HPDF 183K	TUND TUNDOCIAD	A+CODOTA						нРООН11	ጸ	нгрон34	R		HPDOI50R))			HPDQI55R			
		-	•		HPDPX12	T	HPDP Y 83	┰	HPDQC34						нРДОН11		нРDQH34			HPDOISO	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\			нРDQI55			

	639	550 584	264 410	410	269	574	466	337	443	443 694
	10 555	555	4 258	m	3	35	113 466	119	3	3 446
	91%	%08 %08	85% 72%	100%	%86	71%	100% 70%	72%	%26	100%
	emb CAA45124.1	dbj BAA13213.1	emb CAA59350.1	gb AAA36368.1	gb AAC27614.1	emb CAA81022.1	gb AAF22155.1 AF1 33093_10	emb CAA57811.1	emb CAB10847.1	gb AAF13710.1 AF1 99488_1
(PUTATIVE isoform 2) [Homo sapiens]	RNA polymerase II 140 kDa subunit. [Homo sapiens]	similar to putative ATP-dependent RNA helicase K03H1.2 of C.elegans(S41025) [Homo sapiens]	Cctg [Xenopus laevis]	neuroleukin [Homo sapiens]	(AF068227) putative transmembrane protein [Homo sapiens]	ribosomal protein S7 [Homo sapiens]	(AF133093) ARD-1 N- acetyltransferase homologue [Mus musculus]	Histone H3 [Asparagus officinalis]	DNA binding protein [Homo sapiens]	(AF199488) beta-actin [Coturnix coturnix
·	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	1851	1852	1854	1858	1859	1860	1862	1863	1864	1869
	HPDQR20R	HPDQR88R	HPDQS25R	нроотзав	HPDQU81 R	HPDQV07 R	НРDQW39 НРDQW39 R	HPDQX13 R	HPDQY23 R	HPDQZ65R
	HPDQR20	HPDQR88	HPDQS25	нРDQТ32	HPDQU81	нРDQV07	НР DQW39	нроохіз		нРDQZ65

_			Japonical	24 A A B 18676 11	72%	4	240
HPDRA44R	1870	blastx.2	interferon regulatory factor 7 [Mus	golwydiadaeth	45%	362	559
			musculus	11 A A 6790K 11	75%	16	189
HPDRA50R	1871	blastx.2	actin [Diadromus oulchellus]	emb CAA02000.1		1 6	07.6
HPDRF65R	1872	blastx.2	ribosomal protein L8	emb CAA82248.1	28% 95%	787	236
			[Homo sapiens]		20%	505	753
_			To milionia	emblCAA77980.1	94%	20	460
HPDRG73R	1874	blastx.2	Kan [Cams tannuaris]		84%	451	528
HPDRM93	1876	blastx.2	ribosomal protein L37	dbj BAA04888.1	%98	12	302
			Homo sapiens	11:1D A A 0/1988 11	100%	17	307
HPDRO04R	1877	blastx.2	ribosomal protein L3/	apjlezeveteser.			
HPDRP36R	1879	blastx.2	GLI-Krupple related	gb AAA59467.1	37%	36	260
			protein pare saperal		78%	304	372
HPDRP41R	1880	blastx.2	UHX1 protein [Homo	gb AAC50450.1	%88	7	739
			sapiens	1.10 A A 01.409 11	01%	81	290
HPDRQ66R	1881	blastx.2	ribosomal protein [Homo sapiens]	emp[CAA81496.1]	2,1,2		555
HPDRQ84R	1882	blastx.2	Nascent polypeptide	emb CAA56869.1	%6/	t ·	0
			alpha subunit [Homo				
HPDRR71R	1883	blastx.2	(AF081484) alpha-	gb AAC31959.1	98%	16 381	390 422
			Thomasapiens				

299 369		208		260		244	0/7				231		573		222				298	516	522	516	522	522	489	486	522
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%69 %86		100%		%09			0,00				%96	:	%06		%56				64%	%88	20%	20%	48%	48%	25%	20%	44%
gb AAF08293.1 AF1 95094 1		emb CAA37039.1		emb CAA34947.1		emb CAA16159.1					dbj BAA11936.1		gb AAA61279.1		gb AAA36278.1				gb AAB25797.1								
(AF195094) gamma actin-like protein Mus	musculus]	peptidylprolyl isomerase [Homo	sapiens]	leukocyte antigen F	Homo sapiens	(AL021366)	CICKO/21 (.4.1 (FRD	finger protein 2)	(isoform 2) [Homo	sapiens]	HLA-A9HH antigen	[Homo sapiens]	vimentin [Homo	sapiens]	MHC HLA-DR2(non-	Dw2/non-Dw12)a.	glycoprotein beta-chain	[Homo sapiens]	type XVI collagen	alpha 1 chain, alpha 1	(XVI) [human, 1						-
blastx.2		blastx.2		blastx.2		blastx.2					blastx.2		blastx.2		blastx.2				blastx.2							٠	
1885		1886		1888		1890					1891		1893		1895				1897					٠			
HPDRS46R		HPDRS87R	,	HPDRT37R		HPDRU03R					HPDRU37R		HPDRV73R		HPDRW09	X			HPDRY37R								
HPDRS46		HPDRS87		HPDRT37		HPDRU03					HPDRU37		HPDRV73		HPDRW09				HPDRY37								

498	492	504	498	522	516	492	207	522	522	489	522	489	450	522	307	477	498	492	510	304	207	498	522	456	498	492	486	88	88
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310	307	5	7	307	7	319	212	310	7	197	370	212	212	7	7	206	2	S	373	212	7	7	310	310	500	188	2	7	5
43%	41%	64%	28%	41%	62%	43%	%19	40%	54%	44%	45%	30%	31%	28%	28%	48%	28%	25%	43%	45%	25%	26%	38%	32%	20%	38%	27%	24%	21%
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46%	44%	23%	20%	51%	41%	53%	44%	23%	45%	48%	24%	48%	45%	23%	45%	42%	51%	43%	45%	47%	46%	40%	48%	23%	23%	48%	46%	20%	45%
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212	212	203	7	2	2	7	7	S	7	7	209	506	506	506	200	2	14	7	2	S	212	203	212	3	\$. 209	188	188	23
41%	41%	40%	48%	44%	20%	44%	48%	51%	48%	48%	45%	43%	45%	41%	41%	54%	52%	48%	20%	20%	41%	42%	38%	20%	45%	45%	38%	36%	52%
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7	7	212	212	188	14	7	2	188	209	212	212	236	212	500	185	185	23	2	23	203	7	212	209	32	23	3	S	197	2
48%	48%	48%	41%	38%	52%	44%	51%	41%	40%	40%	40%	48%	41%	40%	37%	35%	53%	46%	24%	20%	44%	41%	43%	61%	51%	20%	41%	40%	50%
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5	188	7	2	212	310	209	212	212	212	488	432	7	200	7	2	456	7	485	399	488	488	488	429	491	488	451	188	488	488
36%	38%	45%	45%	38%	47%	45%	37%	45%	41%	41%	38%	44%	20%	45%	37%	52%	47%	20%	35%	72%	53%	53%	45%	27%	40%	41%	39%	24%	63%
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537	650	651	484		036	359	041	820	902		436			326			417	633	338				270				634
496	6	106	720	•	,,,	243	272	92			62			3			94	334	3				61				116
20%	%09	35%	81%		1000	%/%	%00	93%	76%		84%			%66			83%	20%	100%		-		%22		-		%68
	emb CAA70356.1		gb AAD51611.1 AF1	53685_1	11 27 2 4 A CIL 16	dbj BAA02647.1		pirlS16510 MHHUM			pir A44282 A44282			emb CAA47708.1			dbj BAA02633.1		gb AAC39268.1				sp Q64152 BTF3_M	OUSE	•		pir S03957 R5RT18
	Lsc homologue [Homo	sapiens]	(AF153685) truncated	calcium binding protein	[money sapiens]	proteasome subunit	HSIN3 Homo sapiens	Ig mu chain C region,	membrane-bound	splice form - human	retrovirus-related pol	polyprotein pseudogene	- human 1	immunoglobulin M	heavy chain [Homo	sapiens]	APEX nuclease [Homo	sapiens]	(AF067370)	cytoplasmic dynein	light chain; Tctex-1	[Bos taurus]	TRANSCRIPTION	FACTOR BTF3 (RNA	POLYMERASE B	TRANSCRIPTION 1	ribosomal protein L18a - rat
	blastx.2		blastx.2		1.1-1-1.0	DIASTX.2		blastx.2			blastx.2			blastx.2			blastx.2		blastx.2				blastx.2				blastx.2
	1899		1900		,001	1061		1903			1904	_		9061			1908		1909				1911			-	1912
	HPDRZ29R		HPDVA01	<u>-</u> حد	TIPLITY	HPDVA06	K	HPDVB70R			HPDVC28R			HPDVG06	&		HPDV125R		HPDV195R				HPDVK79	&		-	HPDVK93 R
	HPDRZ29		HPDVA01		TIPLITAGE	HPDVA00		HPDVB70			HPDVC28			HPDVG06			HPDVI25		S6IAG4H				HPDVK79				НРDVК93

303	293 625 660 561 546 570	268	573	406			693
4	96 269 418 409 418 326 418	S	_{pre} d	35 4	1 188 396 334	240	163
100%	98% 60% 22% 31% 31%	44%	%16	91% 100%	96% 84% 67% 41%	54% 78%	%96
gb AAB39928.1	emb CAA72789.1	gb AAD48374.1	gb AAF21613.1	gb[AAA36105.1	gb[AAC72274.1	gb AAA39345.1	emb CAA74245.1
cell cycle checkpoint control protein [Homo saniens]	hNop56 [Homo sapiens]	(AF123880) gag polyprotein [multiple sclerosis associated retrovirus element]	(AF019036) human IgG1 neutralizing heavy chain with leader 1	gamma-interferon- inducible protein precursor [Homo sapiens]	(AC005954) ZO-3 [Homo sapiens]	JUN-D protein [Mus musculus]	protein phosphatase 2C
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
1914	1915	1916	1917	1918	1919	1923	1924
15R	HPDVL52R	HPDVM61 R	HPDVM63 R	HPDVM86 HPDVM86 R	HPDVO67 R	HPDVT37R	HPDVU28
HPDVL45 HPDVL	HPDVL52	HPDVM61	НР DVM63	HPDVM86	HPDV067	HPDVT37	HPDVU28

similar to the gb AAA19604.1 100% Drosophila splicing regulator, 1 [Homo 37% regulator, 1 [Homo sapiens] 72% gluathione peroxidase gb AAA59461.1 91% [Homo sapiens] gb AAA59461.1 85% calcium binding protein dbj BAA23325.1 94% [Homo sapiens] emb CAB04787.1 85% GlcNac-1-P transferase emb CAA36054.1 79% (458 AA) [Homo sapiens] 100% ribosomal protein S26 gb AAC26987.1 100% [Homo sapiens] emb CAB06483.1 99% [Homo sapiens] emb CAB06483.1 82% polyadenylate binding gb AAB97309.1 82% protein [Homo sapiens] emb CAA56869.1 91% Nascent polypeptide emb CAA56869.1 91% associated complex alpha subunit [Homo 91%
gb AAA67540.1 gb AAA59461.1 n dbj BAA23325.1 emb CAB04787.1 gb AAC26987.1 emb CAB06483.1 gb AAB97309.1 gb AAB97309.1 emb CAA56869.1
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[Homo saniens]	[emordae ontotr]	n67 mvc protein	THomo sapiens	ribosomal protein S6	[Homo sapiens]	cell surface	glycoprotein [Homo	5S ribosomal protein	[Mus musculus]		(AL049783)	hypothetical protein	HOIIIO Sapieriis	cpn10 protein [Bos	taurus	unknown product	specific to adipose	tissue [Homo sapiens]	alanine	aminotransferase	[Homo sapiens]	(AF033095) testis	enhanced gene	transcript protein	Homo 1	90-kDa heat shock	protein [Sus scrofa]
		blacty 2		blastx.2		blastx.2		blastx 2			blastx.2			blastx.2		blastx.2			blastx.2			blastx.2				blastx.2	
		1043		1944		1945		1048	2		1949			1950		1951			1952			1953				1954	
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-		THE WATER	DFUWUTE -	HPDWN65		HPDWW69	X	0500000000	HFD WOOD		HPDW061			HPDW082		HPDWS61			HPDWT55	: :		HPDWT56				HPDWU47	8

DWU55	HPDWUSS HPDWUSS	1955	blastx.2	protein disulfide	dbj BAA08451.1	94%	1	726
	ጸ			isomerase-related		37%		624
			. :	protein (PDIR) [Homo		27%		636
HPDWU60	HPDWU60	1956	blastx.2	(AF151063) HSPC229	gb AAF36149.1 AF1	93%	176	565
	~			[Homo sapiens]	51063_1	%08	598	657
HPDWU63	HPDWU63	1957	blastx.2	cytokeratin 15 (AA 1 - 456) [Homo saniens]	emb CAA30535.1	100%	2	370
HSCPJ17	HSCPJ17R	1962	blastx.2	(AL031670)	emblCAB43181.1	%16	260	403
				dJ681N20.2 (similar to	-	%88	405	485
50,41	Tropi A 100	1063	110000	ATBOOK CITCHEN, 11ght 1	Martin 4 A 2 4 A 2 1 11	7077	122	300
DSDIA22	חששותפש	1300	Olasta.2	Alrase o Liomo	1.1.50+26-010115	0///	3,	120
				sapiens		47%	238	306
HSDIX73	HSDIX73R	1965	blastx.2	NADH dehydrogenase	dbj BAA07296.1	%98	2	175
				subunit 4L [Homo		100%	178	258
				sapiens]				
HSDZJ21	HSDZJ21R	1968	blastx.2	cytochrome-c oxidase (EC 1.9.3.1) chain I - western lowland 1	pir C59153 C59153	85%		105
HSIFF84	HSIFF84R	1969	blastx.2	(AK002129) unnamed	dbj BAA92096.1	78%	143	9
				protein product [Homo sapiens]				,
HSKJR50	HSKJR50R	1970	blastx.2	translation initiation	gb AAD03467.1	%98	2	337
				factor 3 47 kDa subunit	- -	%86	7	289
				[Homo sapiens]		20%	169	498
	-					46%	11	247
						45%	342	995
HSLHT27	HSLHT27R	1971	blastx.2	URF AGL (NADH	emb CAA24030.1	%08	2	. 139
				dehydrogenase subunit)		%99	2	121

				[Homo saniens]		 		
HSODB16	HSODB16R	1972	blastx.2	cytochrome oxidase	gb AAA20843.1	93%	1	651
	3			subunit II [Homo				
HSPSB24	HSPSB24R	1973	blastx.2	(AF076191) gamma-	gb AAC26520.1	%06	61	732
! }				actin [Trichosurus				
				vulpecula		,000,	,	003
HSPSB70	HSPSB70R	1974	blastx.2	BBC1 [Homo sapiens]	emb CAA45963.1	100%	1	25.
HSPSB74	HSPSB74R	1975	blastx.2	WT1 [Xenopus laevis]	dbj BAA11522.1	%86	-1	791
HSPSB80	HSPSB80R	1976	blastx.2	unc-18homologue	dbj BAA19482.1	92%	7	691
				Homo sapiens				3
HSPSC30	HSPSC30R	1977	blastx.2	cell surface	gb AAA36033.1	91%	124	318
				glycoprotein [Homo		100%	309	374
7900	TY COURT	1070	Llocty 2	(A F1 61 507) HSPC158	oblA AF29122.11AF1	91%	3	338
HSFSE80	HOLDEOOR	12/0	Olasta:2	[Homo saniens]	61507 1	62%	302	299
				[awardan arrarr]	1	54%	477	626
HSPSG03	HSPSG03R	1979	blastx.2	protein phosphatase 2A	emb CAA84414.1	100%	m	191
				65 kDa regulatory				
				subunit, alpha isoform				
				Sus scrota			3	23.0
HSPSG13	HSPSG13R	1980	blastx.2	protein p68 (AA 1-614)	emb CAA33751.1	95%	728	700
				[Homo sapiens]		0,0/	65,	
HSPSG42	HSPSG42R	1981	blastx.2	(AF056490) cAMP-	gb AAC39763.1	%86		924
				specific				
				phosphodiesterase 8A				
				[Homo sapiens]				
HSPSG50	HSPSG50R	1982	blastx.2	(AB036829) skeletal	dbj BAA92340.1	%08	132	278
				Transport and a second				

		645	700		772			404	498	683	821		755		346	749		432	592	632	339	327		385	537	545
		106	629		89			129	367	480	96		. 3		2	099		26	410	561	7	16		2	385	405
		%26	%99		87%			%46	%56	%69	%88		%68		91%	54%		%76	%86	20%	100%	100%		%66	51%	. 47%
		gb AAC31959.1			dbj BAA36501.1			emb CAA32847.1		•	gb AAA60289.1		gb AAA85654.1		gb AAA35531.1	•		dbj BAA05679.1			gb AAA91461.1	oblAAD02337 11		sp P27361 MK03_H	UMAN	•
enriched inositol	phosphatase [Homo sapiens]	(AF081484) alpha-	tubulin isoform 1	Homo sapiens	(AB015610) ribosomal	protein S4X	[Chlorocebus aethiops]	SP-40,40 prepropetide	(AA -22 to 427) [Homo	sapiens]	ribosomal protein S6	[Homo sapiens]	ribosomal protein L5	[Homo sapiens]	medium tumor antigen-	associated 61-kD	protein [Homo sapiens]	BST-2 [Homo sapiens]			UbcH5C [Homo	(AF044671) MM46	[Homo sapiens]	MITOGEN-	ACTIVATED	PROTEIN KINASE 3
		blastx.2			blastx.2		•	blastx.2			blastx.2		blastx.2		blastx.2			blastx.2	,		blastx.2	blastx.2		blastx.2		:
		1983			1984			1985			1986		1987		1988			1989			1990	1991	! !	1992		•
		HSPSG89R			HSPSH39R			HSPSH41R	•		HSPSH49R		HSPSI65R		HSPSJ71R			HSPSJ72R			HSPSQ22R	HSPSO57R		HSPSY67R		٠
		HSPSG89			HSPSH39			HSPSH41			HSPSH49		HSPSI65		HSPSJ71			HSPSJ72			HSPSQ22	HSPSO57		HSPSY67		

				71-111				
				(50.4.1.7)	1. 10000741	/000	ç	283
HSPSZ69	HSPSZ69R	1993	blastx.2	unnamed protein	emb CAB69291.1	%06 80%	285	317
HSPTA57	HSPTA57R	1994	blastx.2	scar protein [Homo	gb AAA36597.1	64%	337	735
				sapiens		100/0	27.0	
HSPTN57	HSPTN57R	1995	blastx.2	(AB015610) ribosomal	dbj BAA36501.1	25%	7 5	/21
				protein S4X		37%	1/8	200
				[Chlorocebus aethiops]		23%	356	904
HSSDM17	HSSDM17	1996	blastx.2	(AĽ137714)	emb CAB70887.1	94%	55	210
	ద			hypothetical protein				
-				[Homo sapiens]			1	
HTLCU84	HTLCU84R	2001	blastx.2	RNA polymerase II	emb CAA87392.1	71%	185	541
				elongation factor-like				
				protein [Homo sapiens]				
HTSHG06	HTSHG06R	2002	blastx.2	URF 2 (NADH	emb CAA24027.1	87%	107	364
				dehydrogenase subunit)		94%		 C)
				[Homo sapiens]				
HULAP70	HULAP70R	2005	blastx.2	cytochrome c oxidase	gb AAA99313.1	100%	251	349
				subunit VIII precursor	•			
				[Homo sapiens]				
HISGA11	HUSGA11	2006	2006 blastx.2	carbonate dehydratase	emb CAA59331.1	63%	220	4/1
	2			[Homo sapiens]		100%	183	218
HIJVFA39	HUVFA39	2009	blastx.2	extensin [Volvox	emb CAA46283.1	34%	460	65 6
	2			carteri]		34%	430	×
HVCAA31	HVCAA31	2010	blastx.2	myosin regulatory light	gb AAA67367.1	%86	103	819
	R			chain [Homo sapiens]			100	100
HVCAA37	HVCAA37	2011	blastx.2	complement factor B	gb AAA16820.1	%%	322	/80
	2			[Homo sapiens]		%88	7	13/
HVCA A 94	HVCAA94	1	2013 blastx.2	F-1-ATPase beta-	gb AAA30395.1	95%	7	435
こくろろう	77.7						•	

643	512	703	644	664	288	794		612			333	524		•		739	=				414	184	430		291	909	758
440	489	128	486	386	133	09		67			- ;	330				68		-			184	119	305	-	61	64	633
20%	100%	%9/	%29	92%	88%	%68		%86			%86	%0/	•	,		%66					%29	%98	38%		72%	%66	47%
		emb CAA66609.1		emb CAA55946.1		dbj BAA02656.1		dbj BAA03401.1			gb AAA52496.1					emb CAB38627.1	-				dbj BAA01569.1				emb CAA55341.1	gb AAA36161.1	
subunit precursor [Bos	taurus]	calcyphosine [Homo	sapiens]	40S ribosomal protein	S12 [Sus scrofa]	human homolog of	DnaJ protein [Homo	'human homologue of	rat ribosomal protein	Ly [HOIDO Sapiens]	glyceraldehyde 3-	phosphate	dehydrogenase (EC	1.2.1.12) [Homo	sapiens]	(AL022721)	dJ109F14.2 (60S	Ribosomal Protein	RPL10A) [Homo	sapiens]	diphtheria toxin	receptor associated	protein [Chlorocebus	aethiops	translin [Homo sapiens]	laminin-binding protein	[Homo sapiens]
		blastx.2		blastx.2		blastx.2		blastx.2			blastx.2					blastx.2					blastx.2				blastx.2	blastx.2	
		2014		2015		2016		2018			2019					2020					2021				2022	2023	
R		HVCAB02	R	HVCAB03	ጸ	HVCAB18	ಜ	HVCAB52	K		HVCAB57	ጸ		·		HVCAB88	R				HVCAC42	&			HVCAD52 R	HVCAE01	ጸ
		HVCAB02		HVCAB03 HVCAB03		HVCAB18		HVCAB52		_	HVCAB57					HVCAB88					HVCAC42 HVCAC42				HVCAD52	HVCAE01 HVCAE01	

:	:	Ŀ		[Homo sapiens]		- - -		
HVCAK02	HVCAK02	2038	blastx.2	hnRNP-E2 [Homo	emb CAA55015.1	%68	96	467
	~			sapiens]		%06	3	92
HVCAL06	HVCAL06	2039	blastx.2	23 kD highly basic	emb CAA40254.1	%96	3	308
	~			protein [Homo sapiens]		30%	305	529
						39%	432	554
HVCA017	HVCAO17 HVCAO17	2040	blastx.2	Nm23 protein [Homo sapiens]	emb CAA35621.1	%02	4	531
HVCA063	HVCA063	2041	blastx.2	similar to tubulin	emb CAA85463.1	%91	82	276
	~			alpha-2 chain		72%	569	322
				[Caenorhabditis	٠.	47%	342	410
		3	,	eregans can		,000	15	3
HVCAP89	HVCAP89R	2042	blastx.2	CIP synthetase	gb[AAB17729.1]	83%	69	512
				homolog [Mus		67%	209	643
				musculus		73%	520	699
,			-			%08	681	710
HVCAQ53	HVCAQ53 R	2043	blastx.2	RPS16 [Homo sapiens]	gb AAA60583.1	100%	.43	480
HVCAR87 HVCAR8	HVCAR87	2045	blastx.2	glyceraldehyde 3-	gb AAA52496.1	74%	54	536
	~			phosphate		-	•	
				dehydrogenase (EC			•	
				1.2.1.12) [Homo				
•				sapiens]				
HVCAS08	HVCAS08R	2046	blastx.2	triose-phosphate	gb AAA35438.1	74%	11	962
				isomerase [Pan		25%	609	803
				troglodytes]				
HVCAS52	HVCAS52R	2047	blastx.2	proteasome subunit C8	dbj BAA00659.1	94%	-	579
				[Homo sapiens]				
HVCAU64	HVCAU64	2049	blastx.2	23 kD highly basic	emb CAA40254.1	%16	57	401

	196	153 107	431	236	636	524	594	478	325	725	629	237	572	535	627	343	647	663	559	256			100	969	1	404	010
	396	215	24	9	274	483	517	452	41	999	290	43	300	353	553	65	327	328	482	23			1	2¢ —		700	404
	%65	76% 90%	%08	85%	41%	71%	45%	%88	%8/	20%	20%	%96	37%	47%	40%	%06	53%	45%	43%	%09				 %66		92%	28%
	dbj BAA91131.1		gb AAA19815.1	-11 A A 74007 11	golada/402/.1				emb CAA40254.1			gb AAD56406.1 AF1	84170 1	1		emb CAA40254.1				dbj BAA06395.1				emb CAA40254.1		gb AAA97405.1	
nrotein [Homo sapiens]	-	product [Homo	in L27	╅	is type	I proteut Litouro	lemondae.		23 kD highly basic	nrotein [Homo sapiens]	ליים ביים ביים ביים ביים ביים ביים ביים	(AF184170) elongation	factor 1-alpha [Sparus	anratal	- Tanana	23 kD highly basic	arotein [Homo caniens]	[aradas arrorr] majord		neurofibromatosis type-	1-GTPase activating-	protein type III [Mus	musculus	23 kD highly basic	protein [Homo sapiens]	AICAR	formyltransferase/IMP
	blastx.2		blastx.2	,	blastx.2				blastx 2			hlasty 2				blocty ?	Olasta:			blastx.2				blastx.2		blastx.2	
	2050		2051		2053				2054	1		2055	203			2056				2057				2058		2060	
0	~		HVCBE76R		HVCBF38R				UTVCBEROR	LIV CDI GOAN		นนุ/กษณา	n v CDCVI	4		111/00021	ונאסט און	-		HVCCA08	8			HVCCK34	2	HVCCV93	R
	HWCRD18		HVCBE76		HVCBF38	_			TIVEDEGO	nvCbroy		1000001					HVCBQ31			HVCCAOR				HVCCK34		HVCCV93	

674	459	341	640	700	267	347	379	326	245	304 383 45
564	61	343	2	215	61	294	59	111	24	32 303 7
%65	%86	95%	%86	39% 40%	39%	38%	47%	91%	87%	98% 88% 61%
	emb CAA40626.1	gb AAB58251.1	emb CAA93540.1	pir A93131 MHDG			gb AAB88484.1	emb CAA04865.1	emb CAB43277.1	emb CAA34066.1
cyclohydrolase bifunctional enzyme [Homo sapiens]	1-8U [Homo sapiens]	similar to mouse Int-6 [Homo sapiens]	3-methyl-adenine DNA glycosylase [Homo sapiens]	Ig mu chain C region - dog			(AF035421) glyceraldehyde 3- phosphate dehydrogenase [Ovis 1	(AJ001612) L-3- phosphoserine- phosphatase homologue [Homo sapiens]	(AL050109) hypothetical protein [Homo sapiens]	ribosomal protein L31 (At 1-125) [Homo sapiens]
	blastx.2	blastx.2.	blastx.2	blastx.2			blastx.2	blastx.2	blastx.2	blastx.2
	2062	2063	2064	2066			2067	2069	2071	2072
	HVCDD19 R	HVCDF50R	HVCDH77 R	HVVAB37 R			HVVAC18 R	HVVAE73 R	HVVAH91 R	HVVAI03R
		HVCDF50	нусрн77	HVVAB37			HVVAC18	HVVAE73	НVVАН91	HVVAI03

CZCU A ATT				The second second		_		
				lambda-chain [Homo				
	7 44 7 44 7	\dashv	1.1.	sapiens)	oblA A 58487.11	%96	3	746
HVVAK46	HVVAK40	//07	DIASLA.2	<u>e</u>		61%	707	169
	4			sapiens		, ,	1	000
HVVAK85	HVVAK85	2079	blastx.2	unnamed protein	emb CAA03726.1	94%	4	3/0
_	R	0000	2.1	product [minestrated]	emblCA A01645.11	81%	157	471
HVVAL81	HVVAL81	7080	blastx.2	phospholipase Az [synthetic construct]			1	
HVVAS27	HVVAS27	2082	blastx.2	ribosomal protein S17	gb AAA60284.1	%66	ა 	413
	RP00B			[Homo sapiens]		2450	,	777
HWVAW2	HVVAW26	2083	blastx.2	23 kD highly basic	emb[CAA40254.1]	/4%	` ;	- 5
9	R			protein [Homo sapiens]		78% 78%	212	503
						2/2/2	6	126
HVVRD91	HVVBD91	2084	blastx.2	gamma non-muscle	emb CAA43140.1	%66	7	430
	RP00B			actin [Oryctolagus			,	
				cuniculus		,000	C	206
HVVBF09	HVVBF09R	2085	blastx.2	(AK001810) unnamed	dbj BAA91922.1	%6%	o	2007
				protein product [Homo			-	
				sapiens		7072	-	610
HVVBH88	HVVBH88	2087	blastx.2	Na,K-ATPase alpha-	dbj BAA00061.1	/4%	-	710
	2			subunit [Homo sapiens]		,86,	,	673
HVVBI16	HVVBI16R	2088	blastx.2	IgG [Homo sapiens]	gb[AAA02914.1]	%69	75	202
						37%	//0	oro
111111111111111111111111111111111111111	4477 478 TO 8 D	2080	hlastx 2	(AB014876) robosomal	dbj BAA34291.1	81%	156	6/4
HVVBJU8	VIOUGA VII	2007	-	protein 1.13 [Cricetulus		%96	40	234
				orisensi		52%	419	60
	1221012	0000	1-10ctv 2	ribosomal protein S17	gb[AAA60284.1]	%66	6	413

	P00B			[Homo sapiens]				
HVVBK13	HVVBK13	2091	blastx.2	(AB022653) anti-	dbj BAA82102.1	%88	73	618
	X			Entamoeba histolytica		62%	633	713
				immunoglobulin kappa		43%	617	733
HVVB065	HVVBO65	2092	blastx.2	immunoglobulin	emblCAA75033.11	%89	12	623
	۲ ک			lambda light chain [Homo sapiens]				
HVVBO88	HVVBO88	2093	blastx.2	immunoglobulin heavy	gb AAA98800.1	82%	77	481
	&			chain VH-III region				
				[Homo sapiens]				
HVVBR70	HVVBR70	2094	blastx.2	immunoglobulin	emb CAA40958.1	%89	20	334
	RP00B			lambda light chain		%19	318	491
				[Homo sapiens]				
HVVBT60	HVVBT60	2096	blastx.2	HYPOTHETICAL	sp Q16465 YZA1_H	100%	448	86
	X			PROTEIN	UMAN			· <u>-</u>
				(FRAGMENT).			_	
HVVCB04	HVVCB04	2098	blastx.2	heavy chain antibody	emb CAA01549.1	%69 ·	99	484
	~			3D6 [synthetic		70%	478	609
				construct]		30%	487	603
IVVCB08	HVVCB08 HVVCB08	2099	blastx.2	(AB022653) anti-	dbj BAA82102.1	%91	73	735
	X			Entamoeba histolytica		_		
				immunoglobulin kappa				
IVVCC06	HVVCC06	2100	hlacty 2	(A BOOS 804) englectin	dhilB A A 31540 11	7007	300	375
)	2000	2	7	(Homo saniens)	1:310100000ion	0/7/	77	2,00
HVVCD81	HVVCD81	2102	blastx.2	glutathione S-	gb[AAA56823.1]	%96	£ 6	640
	R		-	transferase-pi [Homo		2	3	?
			,	sapiens		:		•
	·							

748	214	467	200	3/5	375	375	375	375	378	285	387	378	399	725	349	573	773	540	350		405		
86	7	114				19	19	19	19	_	19	19	28	654	14	310	576	361	e		49		
%2%	94%	84%		95%	%59 %59	65%	%69	63%	52%	63%	47%	44%	44%	70%	%16	%86	%09	25%	100%		100%		
gb AAA36146.1	gb AAA52493.1	emb CAA37883.1		emb CAA73697.1	-		-		.1.0		-				ob/AAC16046.1				gb AAA60078.1		JEN A 25182 11		
nesothelial K7 [Homo	rotein [Homo		Iambda light chanii [Homo sapiens]	FB19 protein [Homo	sapiens]										7 A EO 6103 41 ETD2	Z 11 1 (+C0100-1)	[Homo sapiens]		phosphoglycerate	kinase (EC 2.7.2.3)	(1 DOC 1080) Let 0	(ABU21288) beta 2= microglobulin [Homo	sapiens
blastx.2	blastx.2	blastx.2	7. 31.	blastx.2											C - +	DIASTX.2			blastx.2			blastx.2	
2104	2106	2107		2108											30.0	7109			2110		-+	2111	
HVVCD90 R	HVVCE65	R HVVCF38R		HVVCG29	R											HVVCG31	~		HVVCG46	2		HVVCG93	
HVVCD90	HVVCE65			LIVIVOG29	7700 1 11											HVVCG31			HVVCG46			HVVCG93	

350		119		403	663	739	599	889	261	371		248	307	336	339		583	775	284	378	407	286	87
e		m	,	44	382	644	3	599	1	273		27	242	298	1		14	674	3	283	3	89	4
100%		100%		%68	%62	75%	%96	%08	%46	87%		85%	71%	84%	%66		100%	91%	%56	78%	%89	%68	71%
gb[AAC82471.1]		emb CAA68724.1		emb CAA01550.1			PirlPT0207 PT0207		gb AAB58776.1		•	gb AAA2021.7.1	`		emb CAA77670.1		gb AAA35682.1		gb AAC37563.1		emb CAA25086.1	gb AAA64187.1	
(AF104913) eukaryotic protein synthesis initiation factor [Homo	sapiens]	extracellular matrix	protein Bivi-40 (AA 1 - 303) [Homo sapiens]	3D6 antibody light	chain [synthetic	construct]	Ig gamma chain C	region - chimpanzee	lysophosphatidic acid	acyltransferase-beta	[Homo sapiens]	Ig light chain VII	region [Homo sapiens]		NuMA protein [Homo	sapiens	ribosmal protein small	subunit [Homo sapiens]	Immunoglobulin	lambda chain [Homo	apoferritin H chain [Homo sapiens]	Hin-2 [Homo sapiens]	
blastx.2		blastx.2		blastx.2			blastx.2		blastx.2			blastx.2			blastx.2		blastx.2		blastx.2		blastx.2	blastx.2	
2112		2113		2114			2115		2116			2118			2119		2120		2121		2122	2123	
HVVCH28 HVVCH28 R		HVVCI28R		HVVCI50R			HVVCI85R		HVVCJSIR			HVVCK78	~		HVVCL52	7	HVVCL73	R	HVVCM67 HVVCM67	~	HVVCM84 HVVCM84	HVVCN20	ጸ
HVVCH28		HVVCI28		HVVCI50			HVVCI85		HVVCJ51			HVVCK78 HVVCK78	-		HVVCL52		HVVCL73 HVVCL73		HVVCM67		HVVCM84	HVVCN20 HVVCN20	

337	550 695	346	389 469 501	430 644	302	299	299	290	272	272	299	299	299	272
17	8 549	251	30 359 457	573	3 8 6		9	n	m	m	m	15	15	15
78%	79% 48%	%96	90% 51% 93%	52% 54%	88%	49%	47%	48%	46%	51%	48%	47%	47%	20%
gb AAA73002.1	emb CAA01550.1	gb AAC80170.1	gb AAF12788.1 AF1 91829_1	dbj BAA82518.1	gb AAB69977.1		•			. :				
[Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.], gene product [Homo	3D6 antibody light chain [synthetic	alpha SNAP [Homo	(AF191829) heat-shock protein [Littorina plena]	(AB019120) seven transmembrane receptor [Rattus	(AC002528) alpha2(I) collagen [Homo	sapiens]							•	
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2									
2125	2127	2128	2129	2130	2131									
HVVCN54 R	HVVC013 R	HVVC016	HVVCO87 R	HVVCP41R	HVVCP88R									
HVVCN54	HVVC013	HVVC016	HVVCO87	HVVCP41	HVVCP88									·.

272	302	278	272	272	302	272	314	272	305	536	299	299	302	302	272	299	272	302	272	272	272	278	272	299	299	299	272	302	299
3	m	3	15	n	18	m	18	m	m	n	m	m	m	m	n	m	27	m	m	18	m	15	m	ന	18	18	n	3	3
47%	46%	53%	48%	46%	45%	51%	45%	47%	46%	46%	45%	44%	45%	45%	48%	45%	51%	44%	49%	45%	47%	47%	48%	45%	45%	43%	47%	42%	45%
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272	299	272	272	299	272	302	299	272	272	299	272	272	2000	000	667	302	480	969	629	669	6	507	217		307	
18	15	9	3	18	3	3	c	15	m	ო	3	~	2 "	<u> </u>	<u>.</u>	m	319	583	54	7	.	183	95		65	
48%	45%	42%	44%	43%	48%	42%	43%	44%	43%	43%	43%	76%	40%	0/04	43%	41%	38%	43%	%9 <i>L</i>	7008	2	91% 80%	100%		100%	
							•												gb AAC26520.1	11 4 4 2 7 7 7 11	gb AAA30043.1	emb CAA81626.1	gb AAB69859.1		dbj BAA04889.1	
																			(AF076191) gamma- actin [Trichosurus	vulpecula	90kDa heat shock protein [Homo sapiens]	Rab5c protein [Canis	(AF015283)	Selenoprotein w Home saniens	ubiquitin-like protein	Homo sapiens
																			blastx.2		blastx.2	blastx.2	blastx.2		blastx.2	
	_					,													2133		2135	2136	2137		2141	
																	-		HVVCQ49 R		HVVCQ93	HVVCS28R	HVVCS32R		HVVCU50	α.
										_									HVVCQ49 HVVCQ49 R		HVVCQ93	HVVCS28	HVVCS32		HVVCU50	

292 482 309 479	410	712 747 431 734	715	511	555 716 763	620	477	598	542
309 283 399	es.	71 418 354 654	2	2	1 498 557	8	394	2	423
100% 100% 66% 40%	100%	54% 50% 100% 51%	97%	%66	86% 38% 42%	%06	%96	%86	%56
dbj BAA31199.1	dbj BAA91151.1	gb AAB38382.1	emb CAB58438.1	dbj BAA23366.1	gb AAA49700.1	gb AAD29608.1	emb CAA32574.1	gb AAA36443.1	emb CAA63371.1
(AB010491) natriuretic peptide A type receptor [Homo sapiens]	(AK000419) unnamed protein product [Homo sapiens]	p48 [Homo sapiens]	immunoglobulin gamma-2 heavy chain [Homo sapiens]	Hrs [Homo sapiens]	elongation factor-1 alpha-chain protein (EF-1-alpha) [Xenopus	(AF113887) kappa 1 immunoglobulin light chain [Homo sapiens]	ribosomal protein L34 [Rattus rattus]	phospholipid transfer protein [Homo sapiens]	UDP- GalNAc:polypeptide N-
blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
2142	2144	2146	2147	2148	2149	2150	2151	2152	2153
HVVCV41 R	HVVCW75 R	HVVCX46 R	HVVCY29 R	HVVCY55 R	HVVCY60 R	HVVCY62 R	HVVCY75 R	HVVCY77 R	HVVCZ18 R
	HVVCW7 5	HVVCX46	HVVCY29	HVVCY55	HVVCY60 HVVCY	HVVCY62 HVVCY62	HVVCY75	HVVCY77	HVVCZ18 RVVCZ18

	562	338 502 588	446	155	643 559 683	594	309 593 415	513 485	103
-	20	81 308 541	3	99	8 29 615	4	61 294 287	106 285	2
	85%	80% 74% 62%	%86	%96	95% 25% 56%	%26	92% 39% 48%	64% 56%	%16
	gb AAC39746.1	emb CAA09185.1	gb AAA35507.1	gb AAC15863.1	pir S03894 S03894	emb CAA53625.1	dbj BAA21839.1	dbj BAA14105.1	pir A24815 A24815
acetylgalactosaminyltra nsferase (GalNAc-T3) [Homo sapiens]	(AF013622) immunoglobulin heavy chain variable region [Homo 1	(AJ010446) immunoglobulin kappa light chain [Homo sapiens]	adenylyl cyclase- associated protein [Homo sapiens]	(AF016365) hexokinase I [Homo sapiens]	ADP, ATP carrier protein T2 - human	Lon protease-like protein [Homo sapiens]	SM22 alpha [Homo sapiens]	destrin [Sus scrofa]	calpain (EC 3.4.22.17) large chain 1 - rabbit (fragments)
	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2	blastx.2
	2156	2157	2158	2159	2162	2163	2164	2165	2166
	HVVDH44 R	HVVDM23 R	HVVDM31 R	HVVDM45 R	HVVDN29 R	HVVDN77	HVVDP70 R	HVVDQ46	HVVDQ49 HVVDQ49 R
	HVVDH44	HVVDM2 3	HVVDM3 1	HVVDM4 5	HVVDN29	HVVDN77	HVVDP70	HVVDQ46	нуурQ49

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726	727	746	736	803	612	460		200		229			511	140	140	219	
91	74	069	08	636	121	2		3		7			23	7	n	49	
%68	%56	%89	%88	67%	72%	%86		%62		%98 :			84%	050	% 6 6	%96	,
pir JE0244 JE0244	emb[CAA52808.1]		emb CAA57650.1		gb AAC17119.1	emb CAA63212.1		emb CAA92321.1		emb CAA00676.1	-		gb AAD38038.1 AF1 49822_1	ALIA A B17281 11	go AAB1/381.1	gb[AAB00807.1]	
Ig kappa chain NIG2 precursor - human	gamma subunit of CCT	chaperonin [Homo sapiens]	hevin [Homo sapiens]		(AF065388) tetraspan NET-1 [Homo sapiens]	ch-TOG [Homo	sapiens]	retinoic acid induced	gene c [Homo sapiens]	chimeric monoklonal TSH antibody, gamma	chain [synthetic	construct	(AF149822) mitotic checkpoint protein BUR3 (Mus musculus)	Colorin II [Que corofe]	calpain II [5us scroia]	ZZ:beta-Gal' IgG-	binding fusion protein [unidentified cloning 1
2167 blastx.2	blastx.2		blastx.2		blastx.2	blastx.2		blastx.2		blastx.2			blastx.2	bloom 2	Olastx.2	blastx.2	
2167	2168		2169		2171	2172		2174		2175			2177	2101	1817	2183	
HVVDS12 R	HVVDS21	<u>بر</u>	HVVDS35	R	HVVDT29 R	HVVDT44	~	HVVDW02	K	HVVDW61 R			HVVDX90 HVVDX90 R	LIVIT NATIO	R WLME48	HWMIB35	&
HVVDS12 HVVD	HVVDS21		HVVDS35		HVVDT29	HVVDT44		HVVDW0	7	HVVDW6 1			HVVDX90	עמא המעם	8 8	HWMIB35	8

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	emb CAB55274.1			
	(AL035461)	dJ967N21.3 (novel	protein similar to	predicted 1
	WEF90 2185 blastx.2	•		
	2185			
	HWWEF90	2	<u></u>	
	HWWEF9	0	>	

[0048] Table 2 further characterizes certain encoded polypeptides of the invention, by providing the results of comparisons to protein and protein family databases. The first column provides a unique clone identifier, "Clone ID NO:", corresponding to a cDNA clone disclosed in Table 1. The second column provides the unique contig indentifier, "Contig ID:" which allows correlation with the information in Table 1. The third column provides the sequence identifier, "SEQ ID NO:X", for the contig polynucleotide sequences. The fourth column provides the analysis method by which the homology/identity disclosed in the row was determined. The fifth column provides a description of PFam/NR hits having significant matches identified by each analysis. Column six provides the accession number of the PFam/NR hit disclosed in the fifth column. Column seven, "Score/Percent Identity", provides a quality score or the percent identity, of the hit disclosed in column five. Comparisons were made between polypeptides encoded by polynucleotides of the invention and a non-redundant protein database (herein referred to as "NR"), or a database of protein families (herein referred to as "PFam"), as described below.

The NR database, which comprises the NBRF PIR database, the NCBI [0049] GenPept database, and the SIB SwissProt and TrEMBL databases, was made nonredundant using the computer program nrdb2 (Warren Gish, Washington University in Saint Louis). Each of the polynucleotides shown in Table 1, column 3 (e.g., SEQ ID NO:X or the 'Query' sequence) was used to search against the NR database. The computer program BLASTX was used to compare a 6-frame translation of the Query sequence to the NR database (for information about the BLASTX algorithm please see Altshul et al., J. Mol. Biol. 215:403-410 (1990), and Gish et al., Nat. Genet. 3:266-272 (1993)). A description of the sequence that is most similar to the Query sequence (the highest scoring 'Subject') is shown in column five of Table 2 and the database accession number for that sequence is provided in column six. The highest scoring 'Subject' is reported in Table 2 if (a) the estimated probability that the match occurred by chance alone is less than 1.0e-07, and (b) the match was not to a known repetitive element. BLASTX returns alignments of short polypeptide segments of the Query and Subject sequences which share a high degree of similarity; these segments are known as High-Scoring Segment Pairs or HSPs. Table 2 reports the degree of similarity between the Query and the Subject for each HSP as a

percent identity in Column 7. The percent identity is determined by dividing the number of exact matches between the two aligned sequences in the HSP, dividing by the number of Query amino acids in the HSP and multiplying by 100. The polynucleotides of SEQ ID NO:X which encode the polypeptide sequence that generates an HSP are delineated by columns 8 and 9 of Table 2.

The PFam database, PFam version 5.2, (Sonnhammer et al., Nucl. Acids 100501 Res., 26:320-322, (1998)) consists of a series of multiple sequence alignments; one alignment for each protein family. Each multiple sequence alignment is converted into a probability model called a Hidden Markov Model, or HMM, that represents the positionspecific variation among the sequences that make up the multiple sequence alignment (see, e.g., R. Durbin et al., Biological sequence analysis: probabilistic models of proteins and nucleic acids, Cambridge University Press, 1998 for the theory of HMMs). The program HMMER version 1.8 (Sean Eddy, Washington University in Saint Louis) was used to compare the predicted protein sequence for each Query sequence (SEQ ID NO:Y in Table 1) to each of the HMMs derived from PFam version 5.2. A HMM derived from PFam version 5.2 was said to be a significant match to a polypeptide of the invention if the score returned by HMMER 1.8 was greater than 0.8 times the HMMER 1.8 score obtained with the most distantly related known member of that protein family. The description of the PFam family which shares a significant match with a polypeptide of the invention is listed in column 5 of Table 2, and the database accession number of the PFam hit is provided in column 6. Column 7 provides the score returned by HMMER version 1.8 for the alignment. Columns 8 and 9 delineate the polynucleotides of SEQ ID NO:X which encode the polypeptide sequence which shows a significant match to a PFam protein family.

[0051] As mentioned, columns 8 and 9 in Table 2, "NT From" and "NT To", delineate the polynucleotides of "SEQ ID NO:X" that encode a polypeptide having a significant match to the PFam/NR database as disclosed in the fifth column of Table 2. In one embodiment, the invention provides a protein comprising, or alternatively consisting of, a polypeptide encoded by the polynucleotides of SEQ ID NO:X delineated in columns 8 and 9 of Table 2. Also provided are polynucleotides encoding such proteins, and the complementary strand thereto.

The nucleotide sequence SEQ ID NO:X and the translated SEQ ID NO:Y are sufficiently accurate and otherwise suitable for a variety of uses well known in the art and described further below. For instance, the nucleotide sequences of SEQ ID NO:X are useful for designing nucleic acid hybridization probes that will detect nucleic acid sequences contained in SEQ ID NO:X or the cDNA contained in Clone ID NO:Z. These probes will also hybridize to nucleic acid molecules in biological samples, thereby enabling immediate applications in chromosome mapping, linkage analysis, tissue identification and/or typing, and a variety of forensic and diagnostic methods of the invention. Similarly, polypeptides identified from SEQ ID NO:Y may be used to generate antibodies which bind specifically to these polypeptides, or fragments thereof, and/or to the polypeptides encoded by the cDNA clones identified in, for example, Table 1.

Nevertheless, DNA sequences generated by sequencing reactions can contain sequencing errors. The errors exist as misidentified nucleotides, or as insertions or deletions of nucleotides in the generated DNA sequence. The erroneously inserted or deleted nucleotides cause frame shifts in the reading frames of the predicted amino acid sequence. In these cases, the predicted amino acid sequence diverges from the actual amino acid sequence, even though the generated DNA sequence may be greater than 99.9% identical to the actual DNA sequence (for example, one base insertion or deletion in an open reading frame of over 1000 bases).

Accordingly, for those applications requiring precision in the nucleotide sequence or the amino acid sequence, the present invention provides not only the generated nucleotide sequence identified as SEQ ID NO:X, and a predicted translated amino acid sequence identified as SEQ ID NO:Y, but also a sample of plasmid DNA containing cDNA Clone ID NO:Z (deposited with the ATCC on June 5, 2000 and were given ATCC Deposit Nos. PTA-1982 and PTA-1985; and/or as set forth, for example, in Table 1, 6 and 7). The nucleotide sequence of each deposited clone can readily be determined by sequencing the deposited clone in accordance with known methods. Further, techniques known in the art can be used to verify the nucleotide sequences of SEQ ID NO:X niques known in the art can be used to verify the nucleotide sequences of SEQ ID NO:X.

[0055] The predicted amino acid sequence can then be verified from such deposits. Moreover, the amino acid sequence of the protein encoded by a particular clone can also be directly determined by peptide sequencing or by expressing the protein in a suitable host cell containing the deposited human cDNA, collecting the protein, and determining its sequence.

RACE Protocol For Recovery of Full-Length Genes

Partial cDNA clones can be made full-length by utilizing the rapid [0056] amplification of cDNA ends (RACE) procedure described in Frohman, M.A., et al., Proc. Nat'l. Acad. Sci. USA, 85:8998-9002 (1988). A cDNA clone missing either the 5' or 3' end can be reconstructed to include the absent base pairs extending to the translational start or stop codon, respectively. In some cases, cDNAs are missing the start codon of translation. The following briefly describes a modification of this original 5' RACE procedure. Poly A+ or total RNA is reverse transcribed with Superscript II (Gibco/BRL) and an antisense or complementary primer specific to the cDNA sequence. The primer is removed from the reaction with a Microcon Concentrator (Amicon). The first-strand cDNA is then tailed with dATP and terminal deoxynucleotide transferase (Gibco/BRL). Thus, an anchor sequence is produced which is needed for PCR amplification. The second strand is synthesized from the dA-tail in PCR buffer, Taq DNA polymerase (Perkin-Elmer Cetus), an oligo-dT primer containing three adjacent restriction sites (XhoI, Sall and ClaI) at the 5' end and a primer containing just these restriction sites. This double-stranded cDNA is PCR amplified for 40 cycles with the same primers as well as a nested cDNA-specific antisense primer. The PCR products are size-separated on an ethidium bromide-agarose gel and the region of gel containing cDNA products the predicted size of missing protein-coding DNA is removed. cDNA is purified from the agarose with the Magic PCR Prep kit (Promega), restriction digested with XhoI or Sall, and ligated to a plasmid such as pBluescript SKII (Stratagene) at XhoI and EcoRV sites. This DNA is transformed into bacteria and the plasmid clones sequenced to identify the correct protein-coding inserts. Correct 5' ends are confirmed by comparing this sequence with the putatively identified homologue and overlap with the partial cDNA clone. Similar methods known in the art and/or commercial kits are used to amplify and recover 3' ends.

Several quality-controlled kits are commercially available for purchase. Similar reagents and methods to those above are supplied in kit form from Gibco/BRL for both 5' and 3' RACE for recovery of full length genes. A second kit is available from Clontech which is a modification of a related technique, SLIC (single-stranded ligation to single-stranded cDNA), developed by Dumas et al., Nucleic Acids Res., 19:5227-32 (1991). The major differences in procedure are that the RNA is alkaline hydrolyzed after reverse transcription and RNA ligase is used to join a restriction site-containing anchor primer to the first-strand cDNA. This obviates the necessity for the dA-tailing reaction which results in a polyT stretch that is difficult to sequence past.

[0058] An alternative to generating 5' or 3' cDNA from RNA is to use cDNA library double-stranded DNA. An asymmetric PCR-amplified antisense cDNA strand is synthesized with an antisense cDNA-specific primer and a plasmid-anchored primer. These primers are removed and a symmetric PCR reaction is performed with a nested cDNA-specific antisense primer and the plasmid-anchored primer.

RNA Ligase Protocol For Generating The 5' or 3' End Sequences To Obtain Full Length Genes

Once a gene of interest is identified, several methods are available for the identification of the 5' or 3' portions of the gene which may not be present in the original cDNA plasmid. These methods include, but are not limited to, filter probing, clone enrichment using specific probes and protocols similar and identical to 5' and 3' RACE. While the full length gene may be present in the library and can be identified by probing, a useful method for generating the 5' or 3' end is to use the existing sequence information from the original cDNA to generate the missing information. A method similar to 5' RACE is available for generating the missing 5' end of a desired full-length gene. (This method was published by Fromont-Racine et al., Nucleic Acids Res., 21(7):1683-1684 (1993)). Briefly, a specific RNA oligonucleotide is ligated to the 5' ends of a population of RNA presumably containing full-length gene RNA transcript. A primer set containing a primer specific to the ligated RNA oligonucleotide and a primer specific to a known sequence of the gene of interest, is used to PCR amplify the 5' portion of the desired full length gene which may then be sequenced and used to generate the full length gene. This

method starts with total RNA isolated from the desired source, poly A RNA may be used but is not a prerequisite for this procedure. The RNA preparation may then be treated with phosphatase if necessary to eliminate 5' phosphate groups on degraded or damaged RNA which may interfere with the later RNA ligase step. The phosphatase if used is then inactivated and the RNA is treated with tobacco acid pyrophosphatase in order to remove the cap structure present at the 5' ends of messenger RNAs. This reaction leaves a 5' phosphate group at the 5' end of the cap cleaved RNA which can then be ligated to an RNA oligonucleotide using T4 RNA ligase. This modified RNA preparation can then be used as a template for first strand cDNA synthesis using a gene specific oligonucleotide. The first strand synthesis reaction can then be used as a template for PCR amplification of the desired 5' end using a primer specific to the ligated RNA oligonucleotide and a primer specific to the known sequence of the ovarian antigen of interest. The resultant product is then sequenced and analyzed to confirm that the 5' end sequence belongs to the relevant ovarian antigen.

The present invention also relates to vectors or plasmids, which include [0060] such DNA sequences, as well as the use of the DNA sequences. The material deposited with the ATCC (deposited with the ATCC on June 5, 2000 and were given ATCC Deposit Nos. PTA-1982 and PTA-1985; and/or as set forth, for example, in Table 1, 6 and 7) is a mixture of cDNA clones derived from a variety of human tissue and cloned in either a plasmid vector or a phage vector, as shown, for example, in Table 7. These deposits are referred to as "the deposits" herein. The tissues from which some of the clones were derived are listed in Table 7, and the vector in which the corresponding cDNA is contained is also indicated in Table 7. The deposited material includes cDNA clones corresponding to SEQ ID NO:X described, for example, in Table 1 (Clone ID NO:Z). A clone which is isolatable from the ATCC Deposits by use of a sequence listed as SEQ ID NO:X, may include the entire coding region of a human gene or in other cases such clone may include a substantial portion of the coding region of a human gene. Furthermore, although the sequence listing may in some instances list only a portion of the DNA sequence in a clone included in the ATCC Deposits, it is well within the ability of one skilled in the art to sequence the DNA included in a clone contained in the ATCC Deposits by use of a sequence (or portion thereof) described in, for example Tables 1A or

2 by procedures hereinafter further described, and others apparent to those skilled in the art.

[0061] Also provided in Table 7 is the name of the vector which contains the cDNA clone. Each vector is routinely used in the art. The following additional information is provided for convenience.

[0062] Vectors Lambda Zap (U.S. Patent Nos. 5,128,256 and 5,286,636), Uni-Zap XR (U.S. Patent Nos. 5,128,256 and 5,286,636), Zap Express (U.S. Patent Nos. 5,128,256 and 5,286,636), pBluescript (pBS) (Short, J. M. et al., Nucleic Acids Res. 16:7583-7600 (1988); Alting-Mees, M. A. and Short, J. M., Nucleic Acids Res. 17:9494 (1989)) and pBK (Alting-Mees, M. A. et al., Strategies 5:58-61 (1992)) are commercially available from Stratagene Cloning Systems, Inc., 11011 N. Torrey Pines Road, La Jolla, CA, 92037. pBS contains an ampicillin resistance gene and pBK contains a neomycin resistance gene. Phagemid pBS may be excised from the Lambda Zap and Uni-Zap XR vectors, and phagemid pBK may be excised from the Zap Express vector. Both phagemids may be transformed into E. coli strain XL-1 Blue, also available from Stratagene.

[0063] Vectors pSport1, pCMVSport 1.0, pCMVSport 2.0 and pCMVSport 3.0, were obtained from Life Technologies, Inc., P. O. Box 6009, Gaithersburg, MD 20897. All Sport vectors contain an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, also available from Life Technologies. See, for instance, Gruber, C. E., et al., *Focus 15:59-* (1993). Vector lafmid BA (Bento Soares, Columbia University, New York, NY) contains an ampicillin resistance gene and can be transformed into *E. coli* strain XL-1 Blue. Vector pCR[®]2.1, which is available from Invitrogen, 1600 Faraday Avenue, Carlsbad, CA 92008, contains an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, available from Life Technologies. See, for instance, Clark, J. M., Nuc. Acids Res. 16:9677-9686 (1988) and Mead, D. et al., Bio/Technology 9: (1991).

[0064] The present invention also relates to the genes corresponding to SEQ ID NO:X, SEQ ID NO:Y, and/or the deposited clone (Clone ID NO:Z). The corresponding gene can be isolated in accordance with known methods using the sequence information disclosed herein. Such methods include preparing probes or primers from the disclosed

sequence and identifying or amplifying the corresponding gene from appropriate sources of genomic material.

Also provided in the present invention are allelic variants, orthologs, and/or species homologs. Procedures known in the art can be used to obtain full-length genes, allelic variants, splice variants, full-length coding portions, orthologs, and/or species homologs of ovarian associated genes corresponding to SEQ ID NO:X or the complement thereof, polypeptides encoded by SEQ ID NO:X or the complement thereof, and/or the cDNA contained in Clone ID NO:Z, using information from the sequences disclosed herein or the clones deposited with the ATCC. For example, allelic variants and/or species homologs may be isolated and identified by making suitable probes or primers from the sequences provided herein and screening a suitable nucleic acid source for allelic variants and/or the desired homologue.

[0066] The polypeptides of the invention can be prepared in any suitable manner. Such polypeptides include isolated naturally occurring polypeptides, recombinantly produced polypeptides, synthetically produced polypeptides, or polypeptides produced by a combination of these methods. Means for preparing such polypeptides are well understood in the art.

[0067] The polypeptides may be in the form of the secreted protein, including the mature form, or may be a part of a larger protein, such as a fusion protein (see below). It is often advantageous to include an additional amino acid sequence which contains secretory or leader sequences, pro-sequences, sequences which aid in purification, such as multiple histidine residues, or an additional sequence for stability during recombinant production.

[0068] The polypeptides of the present invention are preferably provided in an isolated form, and preferably are substantially purified. A recombinantly produced version of a polypeptide, including the secreted polypeptide, can be substantially purified using techniques described herein or otherwise known in the art, such as, for example, by the one-step method described in Smith and Johnson, Gene 67:31-40 (1988). Polypeptides of the invention also can be purified from natural, synthetic or recombinant sources using techniques described herein or otherwise known in the art, such as, for

example, antibodies of the invention raised against the ovarian polypeptides of the present invention in methods which are well known in the art.

[0069] The present invention provides a polynucleotide comprising, or alternatively consisting of, the nucleic acid sequence of SEQ ID NO:X, and/or the cDNA sequence contained in Clone ID NO:Z. The present invention also provides a polypeptide comprising, or alternatively, consisting of, the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X or a complement thereof, and/or a polypeptide encoded by the cDNA contained in Clone ID NO:Z. Polynucleotides encoding a polypeptide comprising, or alternatively consisting of the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X, and/or a polypeptide encoded by the cDNA contained in Clone ID NO:Z are also encompassed by the invention. The present invention further encompasses a polynucleotide comprising, or alternatively consisting of, the complement of the nucleic acid sequence of SEQ ID NO:X, a nucleic acid sequence encoding a polypeptide encoded by the complement of the nucleic acid sequence of SEQ ID NO:X, and/or the cDNA contained in Clone ID NO:Z.

[0070] Many polynucleotide sequences, such as EST sequences, are publicly available and accessible through sequence databases and may have been publicly available prior to conception of the present invention. Preferably, such related polynucleotides are specifically excluded from the scope of the present invention. Accordingly, for each contig sequence (SEQ ID NO:X) listed in the third column of Table 1, preferably excluded are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 and the final nucleotide minus 15 of SEQ ID NO:X, b is an integer of 15 to the final nucleotide of SEQ ID NO:X, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:X, and where b is greater than or equal to a + 14. More specifically, preferably excluded are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a and b are integers as defined in columns 4 and 5, respectively, of Table 3. In specific embodiments, the polynucleotides of the invention do not consist of at least one, two, three, four, five, ten, or more of the specific polynucleotide sequences referenced by the Genbank Accession No. as disclosed in column 6 of Table 3. In further embodiments, preferably excluded from the invention are the specific polynucleotide

sequence(s) contained in the clones corresponding to at least one, two, three, four, five, ten, or more of the available material having the accession numbers identified in the sixth column of this Table. In no way is this listing meant to encompass all of the sequences which may be excluded by the general formula, it is just a representative example. All references available through these accessions are hereby incorporated by reference in their entirety.

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HDABW50	191	984168	1.510	15 - 524	AW160956, BG112870, AA181146, AL533095, BE299316, BF206076, AL518638, BG025731,

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	15 - 1047	15 - 900	15 - 670
	1 - 1033	1 - 886	1 - 656
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	193	194	195
	HMVAW42	HAGDF03	HOPKU29

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985580	986078	986158
196	197	198
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HNFJH73	199	986165	1 - 2951	15 - 2965	AL135307, AV705200, BG113555, BE893152, AU140178, BE275933, BE894324, AI934515, AA716409, BF060791, BE299687, AI677871, AW954275, N29134, AI808415, AI621031, AA582436, AI923237, BG179963, AA700347, AA463830, AA825773, BE393064, BF219349, AU158345, AI831482, BE893515, AI829698, AI354807, AI041945, BE208699, AA831207, H14671, AA906989, AA460265, N29562, AA679105, H07010, N29294, AA664218, BF438401, N26261, N20226, N21171, K66676, AI147747, BE567629, R53048, A1969497, AA953469, AA976405, AW206458, AW373029, AW503770, BE407841, AA224344, BF11102, ZA4255, AA878064, H54597, Z44642, BG006488, AA303163, H05459, N42173, AI206338, AA219130, AA503077, T29413, BE551030, AI805791, R53135, H54374, BF110041, AI970159, R67394, AA918289, BE3140879, A13097, AA696999, H60931, BF212051, F01802, F03399, A156718, BF976822, H10504, AA137020, AW590683, AI652433, F03681, BF571534, C15517, F05530, AA136924, AA369755, T49871, AA341117, H60845, W93139, AA010506, BF344411, H67688, AI253282, AI825058, T49872, AW798336, AI185461, AA026121, BF701801, AA026205, AA722157, H14581, AA301189, AW411286, BE220882, AX004174, AF075706, AF075706, B06025, B06024, D12765, U35622, U18018, X63190, AF095890, X96377, and X96376.
HNTCH03	200	986328	1 - 738	15 - 752	AI631653, AW514704, AI683239, AI762458, BE544805, AI952420, AI675981, AI926932, BE536677, H48459, AA780363, R99152, AI049877, AA626759, AL031672, D80008, AP001712, AP000466, AL009051, AC005156, AC006006, AC002115, AC008745, AL121825, U91323, AC025030, AL133244, AL161938, AC007883, AP001630, AC016612, AP001748, AC005531, AC004386, AC002418, AL031597, AC005840, AL159970, AC008747, AL080242, AC002978, and AL031651.
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		,			W58692, BE926335, BE544664, R54752, AI183902, AA302461, AI146843, BF760392,
				, ,	AA464316, U66895, L76200, and A11042.
HODEC78	379	100829	1 - 1457	15 - 1471	AA287686, AA207154, AU154168, AA179206, BF959881, A1640238, AU132680, BF194738, DE64644, AW126818, BE77266, AT162680, AT64648, AT673667, AW1264602, AW166551
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					A1346434 RE777693 RENAI364 RE442169 AW013899 HX3373 AA018189 RE276307
					AWGGIDS AA749310 BF433845 AA366409 AA769579 H87151 H87035 A1868789
					A4464267 AA249429 AIR16088 AA471320 BE256470 AW367751 AW391782 BF931655
	•				AI580257, AA101455, BF961680, BE745650, AA973114, U90028, and U90029.
HODEF29	380	100830	1 - 413	15 - 427	AW949656, AW949631, AW949643, AW949618, AW949642, AW949655, D80022, D80212,
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AV718489, AW973307, AW966534, AV718931, AV719557, AV720731, AW959202,
AV699447, AW938993, AV/22801, AW939130, D30263, AW900022, C15077, AW735277, AW96374, AW97387, D80164, AW973541,
AW966054 AW978648, AW966050, D59787, AV719783, D59467, AV719468, AV718800,
AW965185, AW965197, D59275, AW965196, AW965184, AV720791, AV720464, AV718770,
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AW959062, AW964477, AW956434, AW964488, AW949641, AW962082, AV699927,
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 AV701335, AV701166, AV742430, AV701149, AA305578, AV701043, AV701332,
AV701017, AV701248, D80134, AV742048, AV701431, AV699866, AW964468, D80248,
AV720150, AW973490, AW966386, AV645389, AV645344, AW964532, AW966397, D81026,
D80949, AV701130, AW973465, AV701419, AW378532, AW964766, AV701154, AV701422,
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AV720616, AW966049, D51250, AW9511169, AW949620, D80133, AW504033, AV 10123,
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					BE894405, AW387565, AA683545, AA514682, AI912084, BE707577, AA284269, BE892060,
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					AI539420, AA609176, AI803042, AI688409, N36974, AI095658, AA625154, AA454670,
					AI015532, AA532391, AI801045, AW304056, AI078810, AW515831, AA313278, AI143830,
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					AW576279, BF849416, AI225071, N69026, BF922788, AV686309, AW866508, Z46156,
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					AI300698, AA320833, AA994140, AW382298, N39669, W72123, AI183682, AI678525,

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			383	384
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AF019386. AF019385. AF177430, and AJ388534.	BE002704, BE065136, AW152446, AV734696, T62861, AA58449, AW370854, AW295305, AI539492, AI093863, T62711, AA663693, AA808792, AW371354, BE138955, AI343194, AI251970, M18740, M25718, and U13369.	AU121386, BG024628, AL523997, BF794753, AU137819, BG177391, BG1706053, BG229069, BG106999, BG107310, BE265326, BF305474, BF347603, BF796967, BE018909, BE018889, BF797739, BE167696, BF981741, AL039474, BF350360, BE244700, BF869380, AA651690, BF183251, BE778225, BE740869, BE271603, AW474216, D58450, AU124160, AW361236, AW466946, BG107057, BF772633, W27829, AU076945, W26115, BF759430, AW664187, AA570434, BF675704, L12579, M74099, U68542, L43505, AB041595, AC005103, AF047825, AC005086, AF004225, X75013, AC006457, AC002978, and U25750.	BG250587, AV734294, AW949920, AU125757, BF529108, AW952/796, AA4443350, BG250587, AV734294, AW949920, AU125757, BF529108, AW952/796, AA44336, BE673024, AI335961, AA738061, AA43433, BE938159, AW513187, AI860346, AW0099354, BE294816, AU149357, AW058419, BE042809, BG150358, W40461, N95557, AA418263, AA418366, AI183756, BF983116, N52313, AA620306, AA884462, AI001006, D63258, BE274198, AI400765, N93959, BF886486, AW262034, AW183978, BF765839, AU138365, AA249125, BF509213, BF886848, BF887480, AI658872, AI869768, R88859, BF223596, R47440, R47819, BE296217, BF755474, BE707786, AA448501, W24950, AI701222, AA091617, AA704660, BF692008, AW138128, A46722, D85510, D86331, E16655, and AF144759.	BF508974, AW965115, AI630839, AI741643, AW072035, AW268897, AV723999, Docooo, AW072761, AI016072, BF512461, AA344181, C16744, D62943, BF437674, AA344606, AA905880, D62570, AW965114, W60522, F05944, AI915081, AA307598, AV754716, AL042630, AA689351, BF448904, AA429481, AW392414, AI124706, BF883138, AA603400, AW732205, BF769787, AA610381, AL137946, AI471455, AA550959, AW502873, F36200, BF766147, AA132912, AV728369, AI687343, BF742657, H86221, N91711, AI624379, AA228778, AA642809, AV764406, AA828153, AA946848, AA084230, AA578832, T93109, AA52800, AI356440, BE868437, AW949694, BG026818, BG026893, C14480, AA508036, AI754037, AA525293, AI568659, AA610826, AI185394, AV743799, AA362349, AA3624778, AA484892, AA678950, AA135761, W39287, AU150583, AA654778, T05354, AA648922, AA58615, AA468923, BF737749, AA83731, AW511778, AA525331, AV764259, AA728861, F26072, AA445951, AW850230, BF913900, AA4833034, AW151247, BE143664, AA533054, AW979295, AA371410, N71619, AA483483, AA832016, AA831801, AV762430, AI246996, AA54682, AA58863, AA58863, AA58863, AA58863, AA58863, AA58863, AA58863, AA58863, AA831410, N71619, AA483483, AA832016, AA831801, AV762430, AI24996, AA54082, AA58863, AA58863, AA58863, AA58863, AA58863, AA58863, AA831810, N71619, AA483483, AA832016,
4	15 - 783 B	15 - 793 A B B B B B B B B B B B B B B B B B B	15 - 2046	15 - 931
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	385	386	387	
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	1 - 1291	1 - 516	1 - 461	1 - 809
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	389	330	391	392
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	1 - 1705	1 - 363
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	393	394
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	15 - 1497	15-350	15 - 1808
	1 - 1483	1 - 336	1 - 1794
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	408	409	410
	HEZES17	HELHM06	HE2AG50

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					A78862, D34614, AX020190, A30438, D88547, A94995, AR008443, AR060385, AB002449, Y17187, AR025207, I50126, I50132, I50128, I50133, AR008277, AR008281, AR066488,
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					AJ287395, Y09669, A43192, A43190, AR038669, AR054175, AR066487, AR074136, U46128,
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HE2ES61	411	100938	1 - 1258	15 - 1272	AF075305, AW439817, AW439844, AI493164, AA203275, BF690753, AA533476, N31463,
		∞		-	AI435348, AA838687, AI334356, AI282866, AI823812, AV/19236, AA3U/020, AW1388/3, AW334781 AW994877 RF445452 RF673101, AA035519, AW118716, AA219612.
					AW194890, AA737528, AA453621, AC005034, and AB026911.
HE2S043	412	100939	1-1107	15 - 1121	BF345442, BE745400, BF002361, BF591185, AW006311, AI740645, AW972577, AW190738,
		∞			AW872945, BF590839, AW515791, AI337338, AI623327, AI160778, BG052059, A1146339, A1684182 AW130008 AT380309 AT375011 BG222460 AI539558 AI217110, AI371688.
					AW615545, A1360696, AW004738, AA885332, AA506508, AI264485, AI356161, AA280731,
					AW873997, A1479723, A1471972, A1077764, A1356162, A1472175, AA557465, AW007779,
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					AW248081, 123526, A442651, H1788, B4454514, AA8/1037, NOOROUS TATACASE TATA
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HLHC107	413	100940	1 - 517	15 - 531	W31456, W52407, AC005410, AC020744, AL445249, AL132821, AL035652, and AC016925.
HTTBR65	414	100941	1 - 461	15 - 475	BE314959, AW411287, AW411286, BF304101, BF976822, AW411455, BF344411, BF339991,
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		4			BF571534, AW367868, BF809505, W93139, AA010506, AF095890, D12765, E06025,
HPFCV71	415	100945	1-661	15 - 675	AW028647, R31337, BF892076, BF892569, AW816131, BF988799, BF993589, BF990226,
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			-		AW080772, AW157173, AA878955, AW973825, BE246411, BG260565, AA932787,
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					AC005695, AL138960, and AL122014.
HCABR46	416	100947	1 - 869	15 - 883	AW131143, BF343353, AW166982, AI922723, AA769108, AW084960, AI076939,
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